

FIG. 1

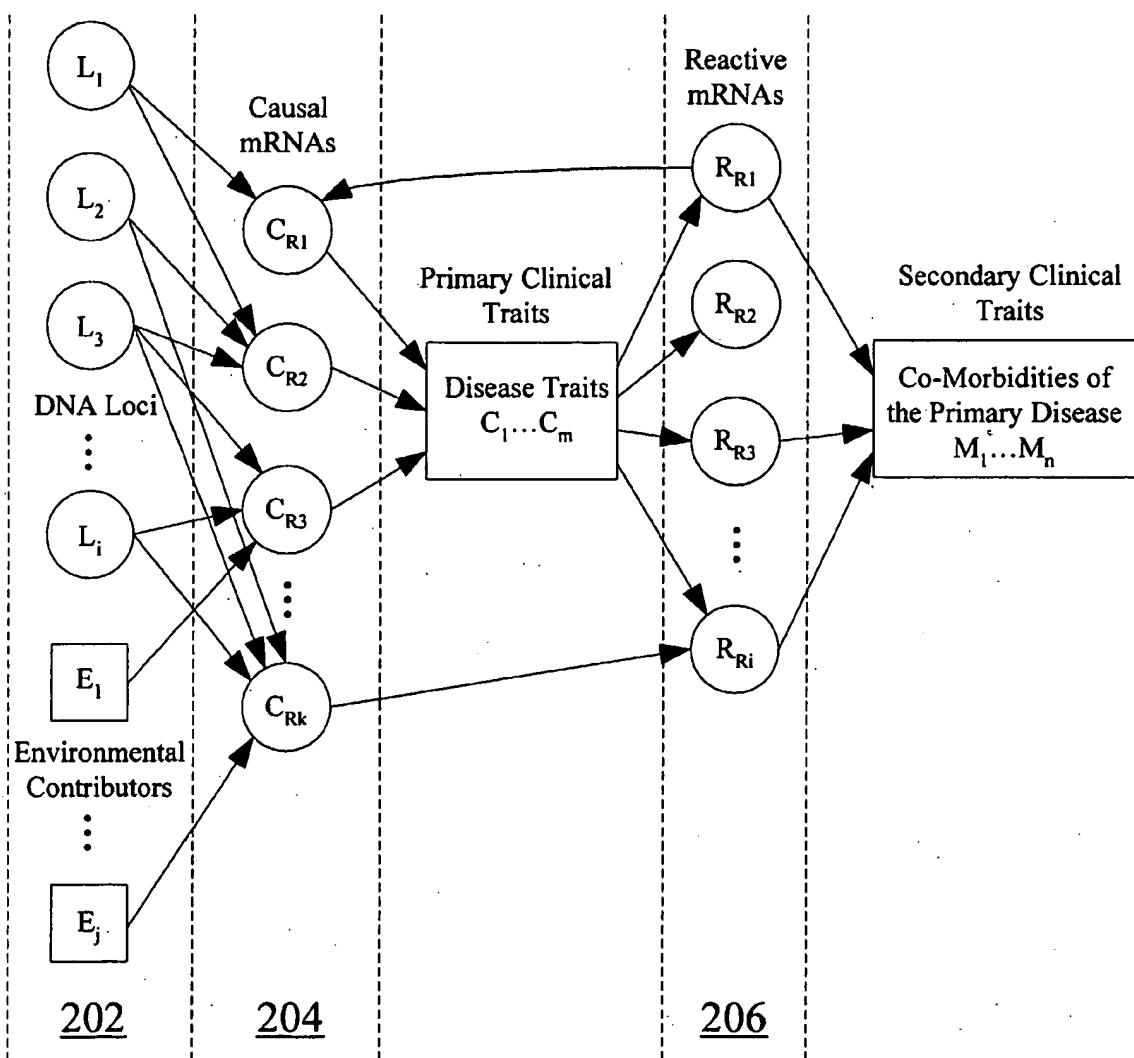
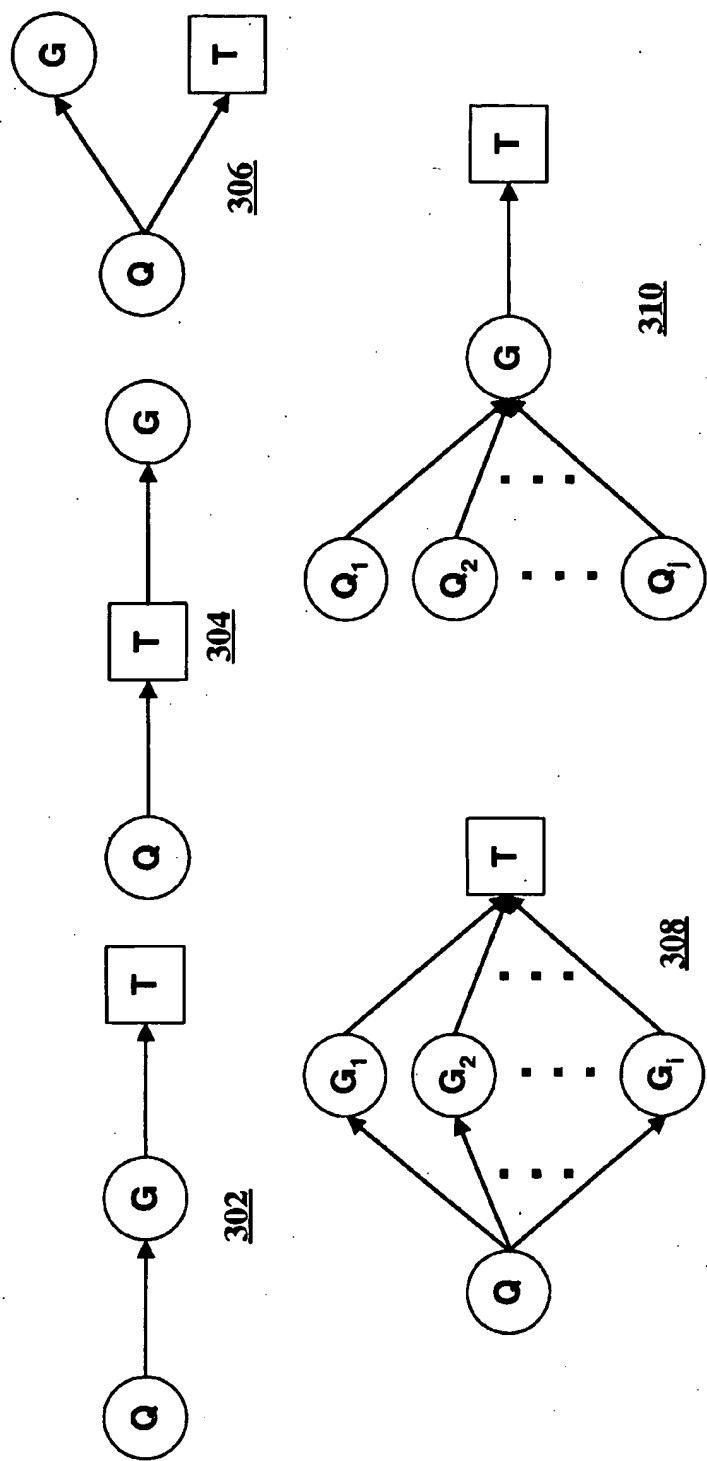


Fig. 2

Fig 3A  
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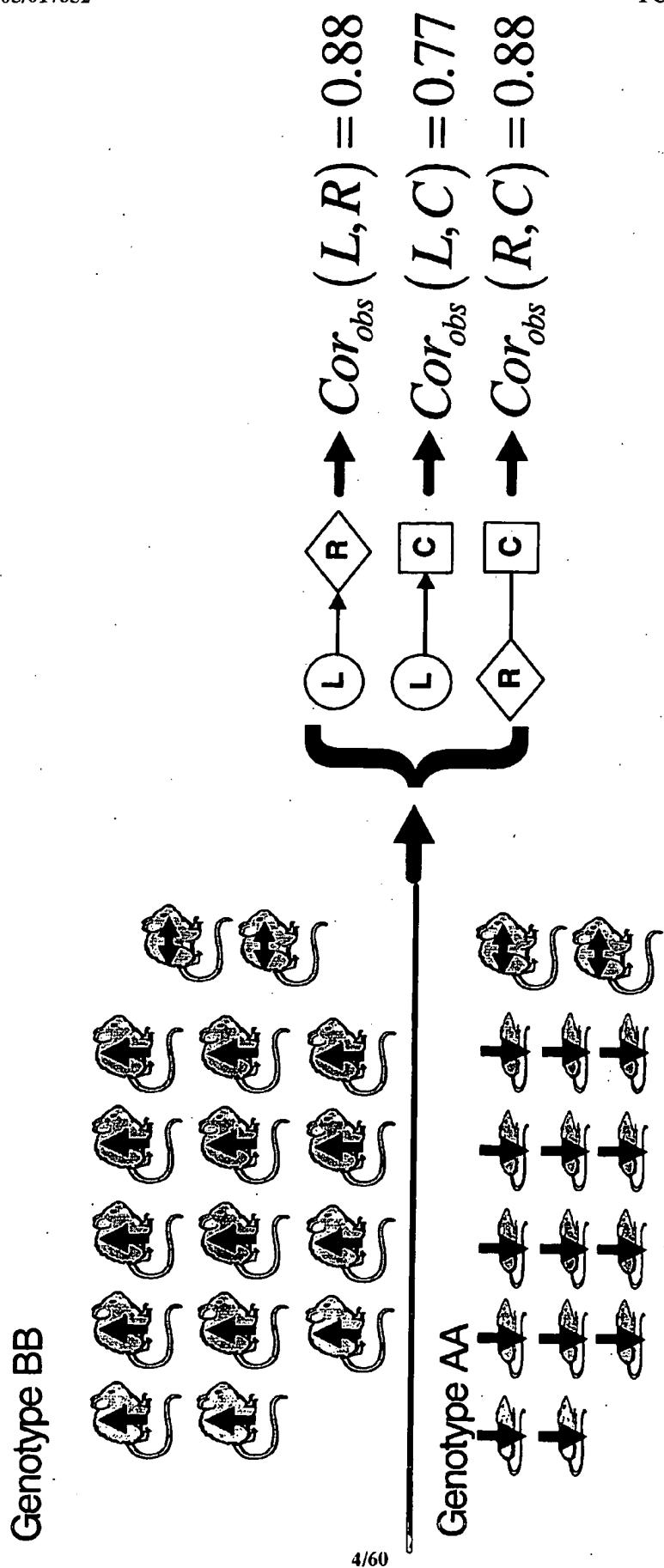


Figure 3B

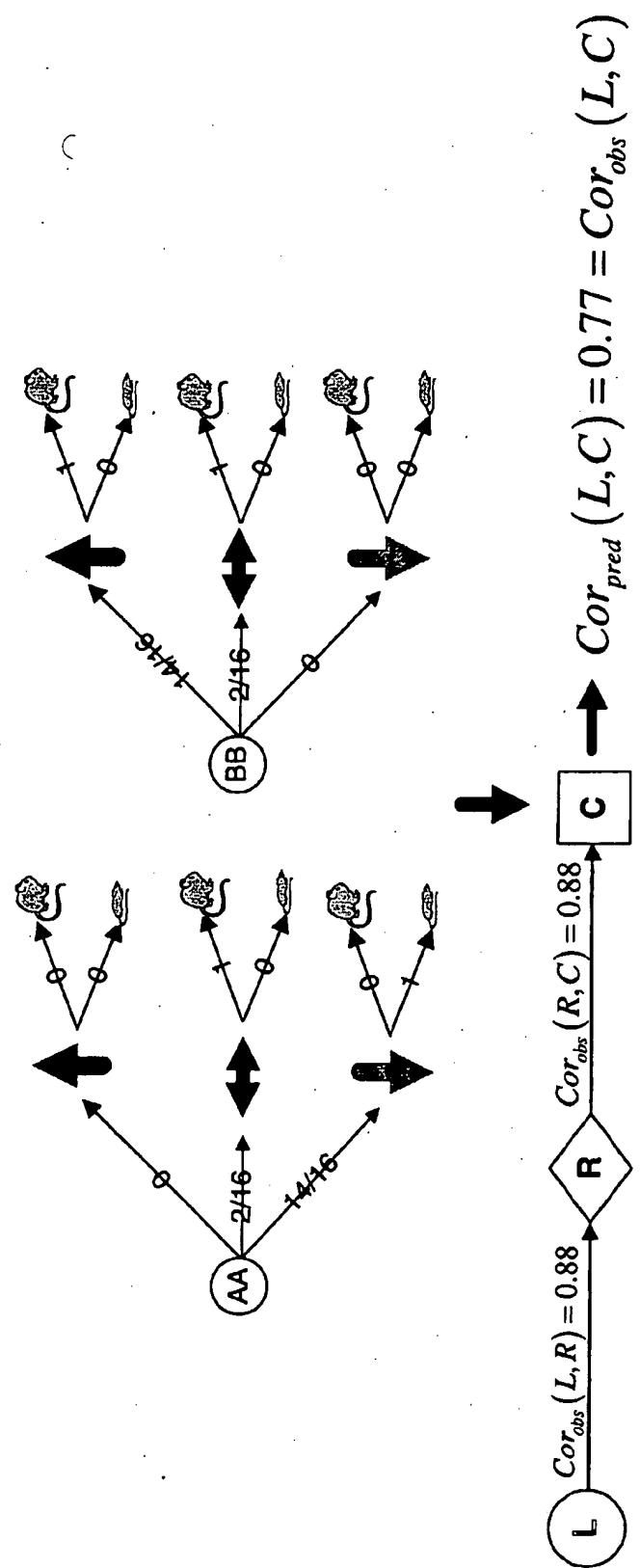


Figure 3C

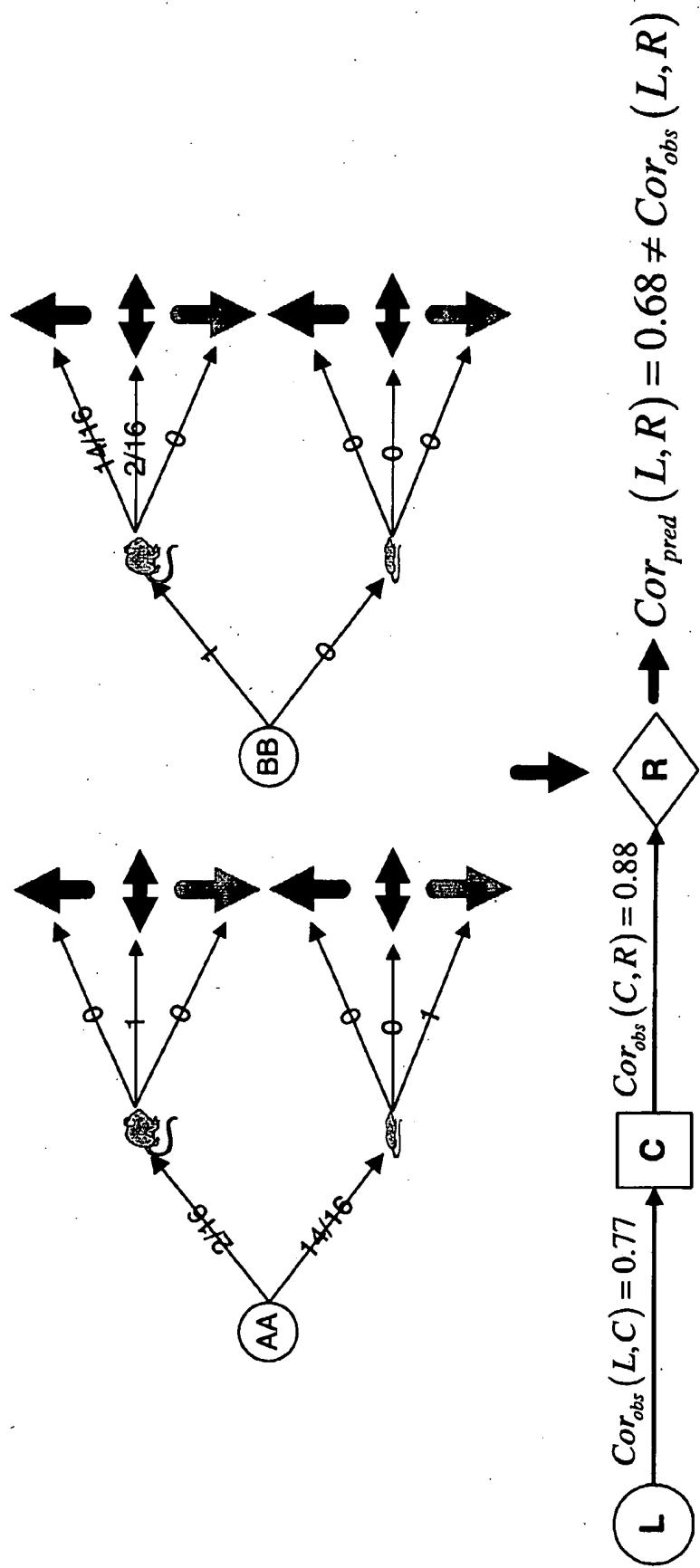


Figure 3D

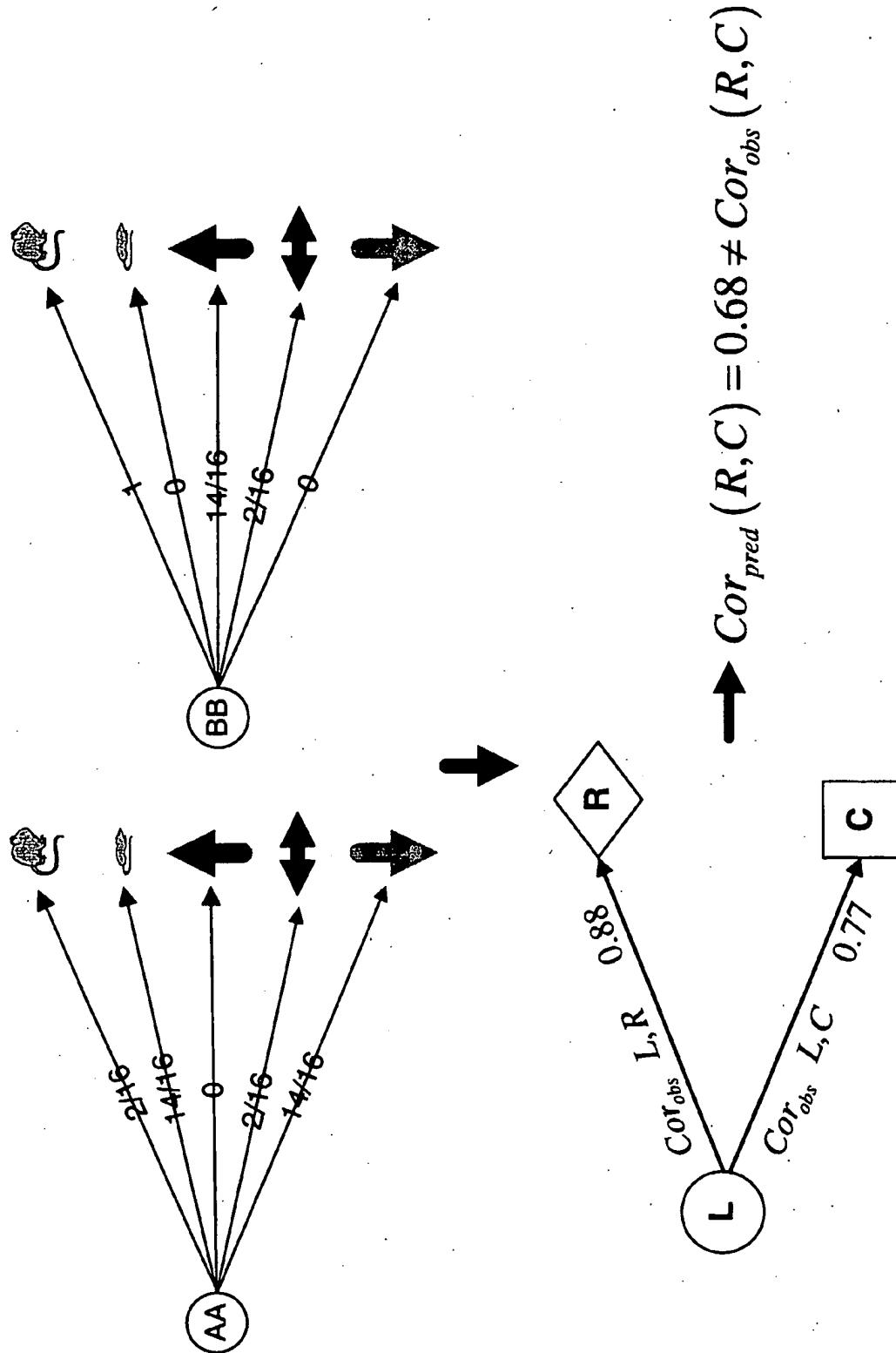


Figure 3E

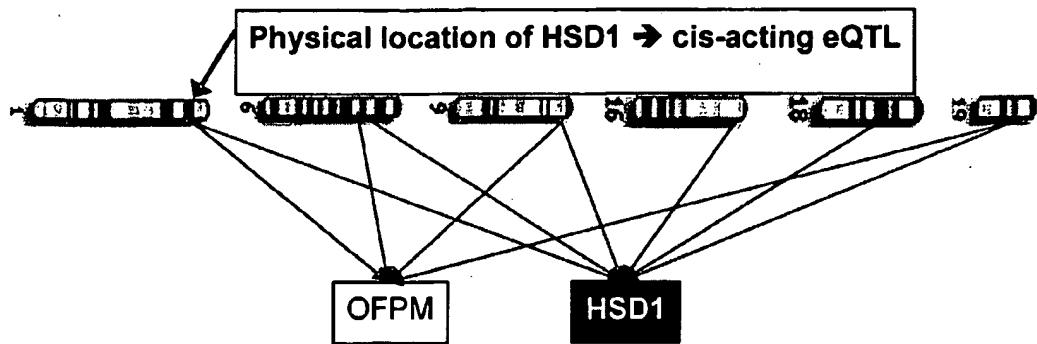


Fig. 4.

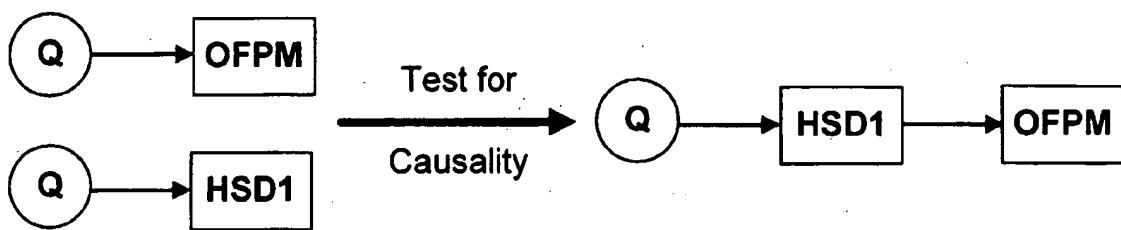


Fig. 5

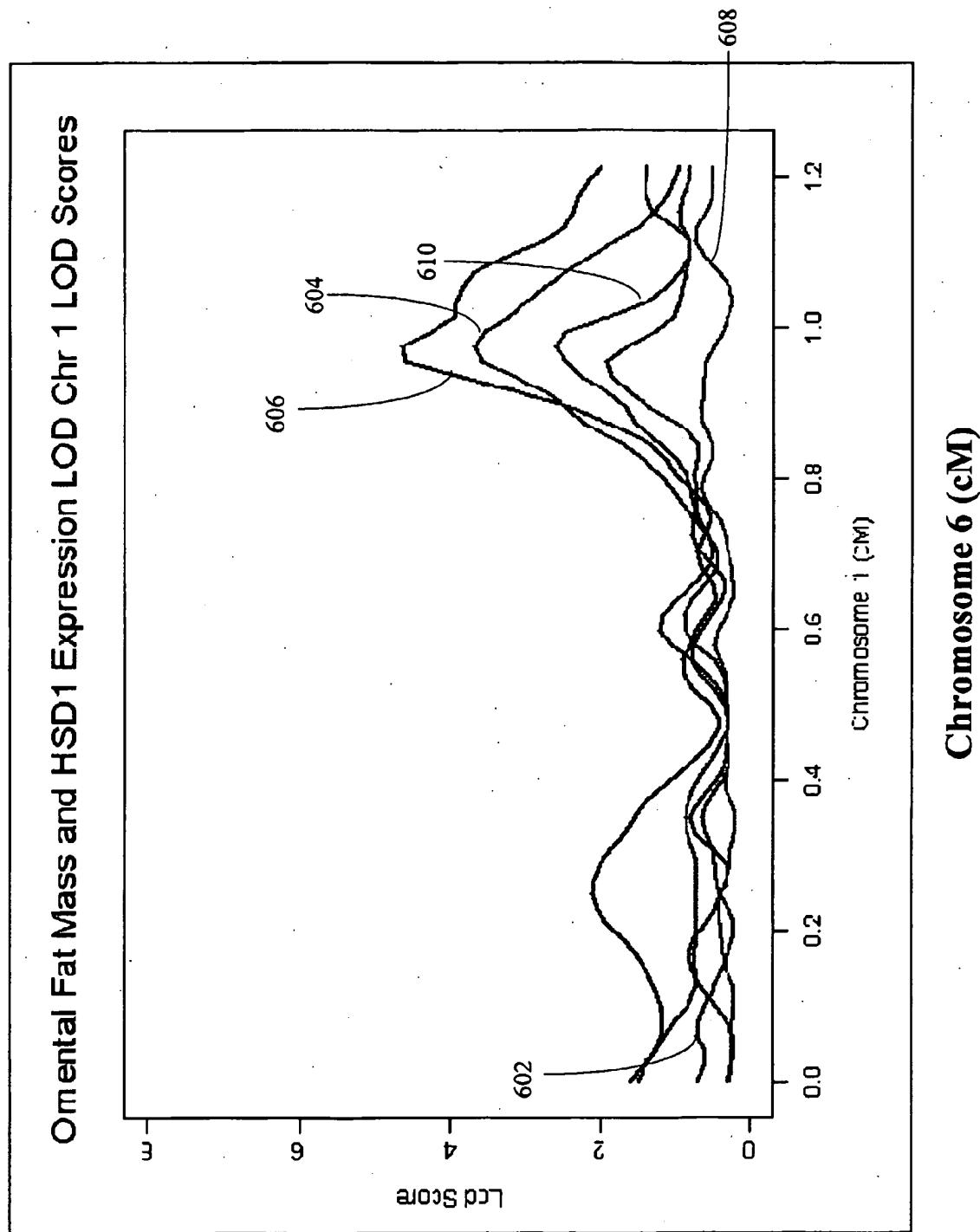


Fig. 6

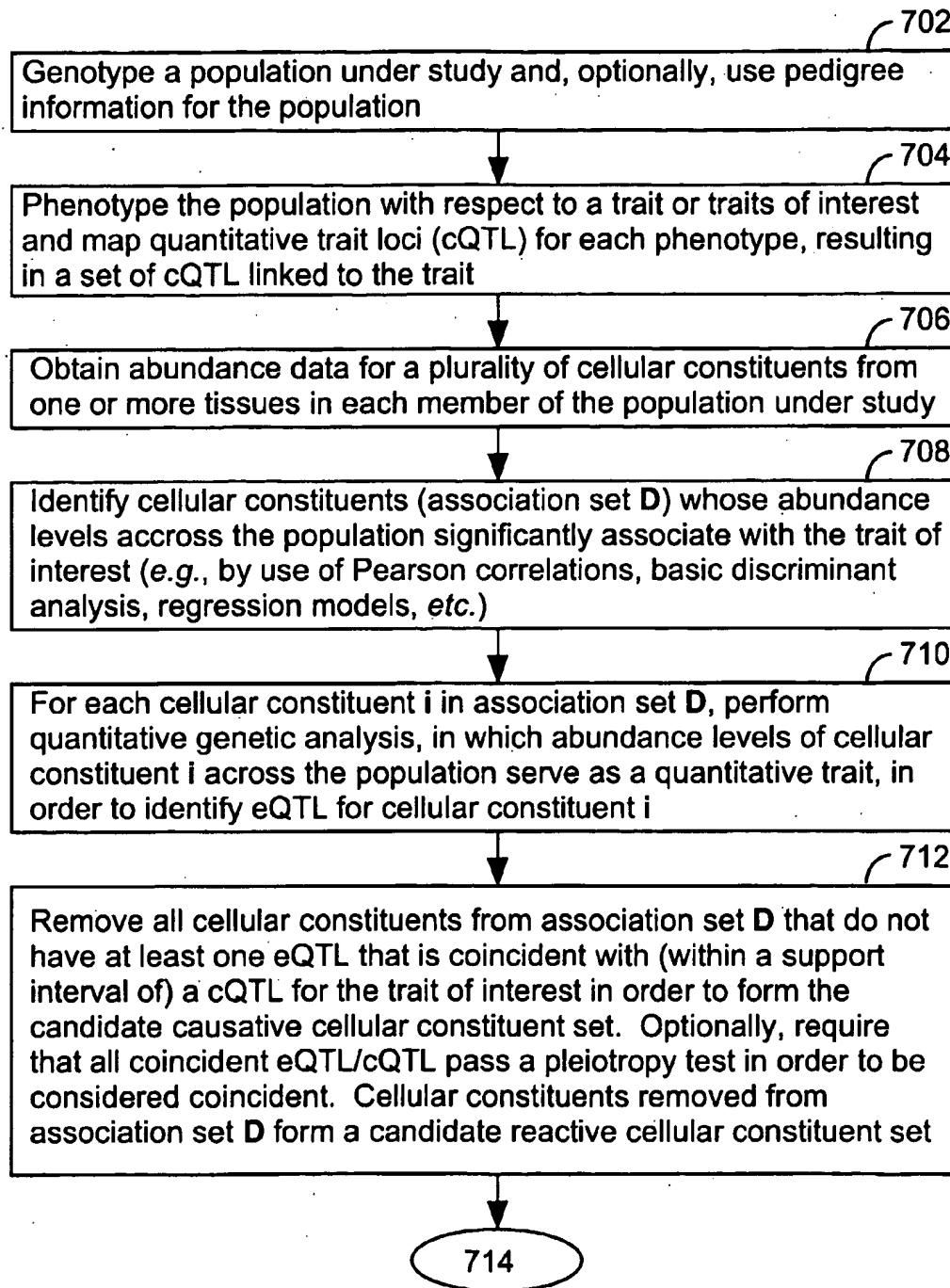


FIG. 7A

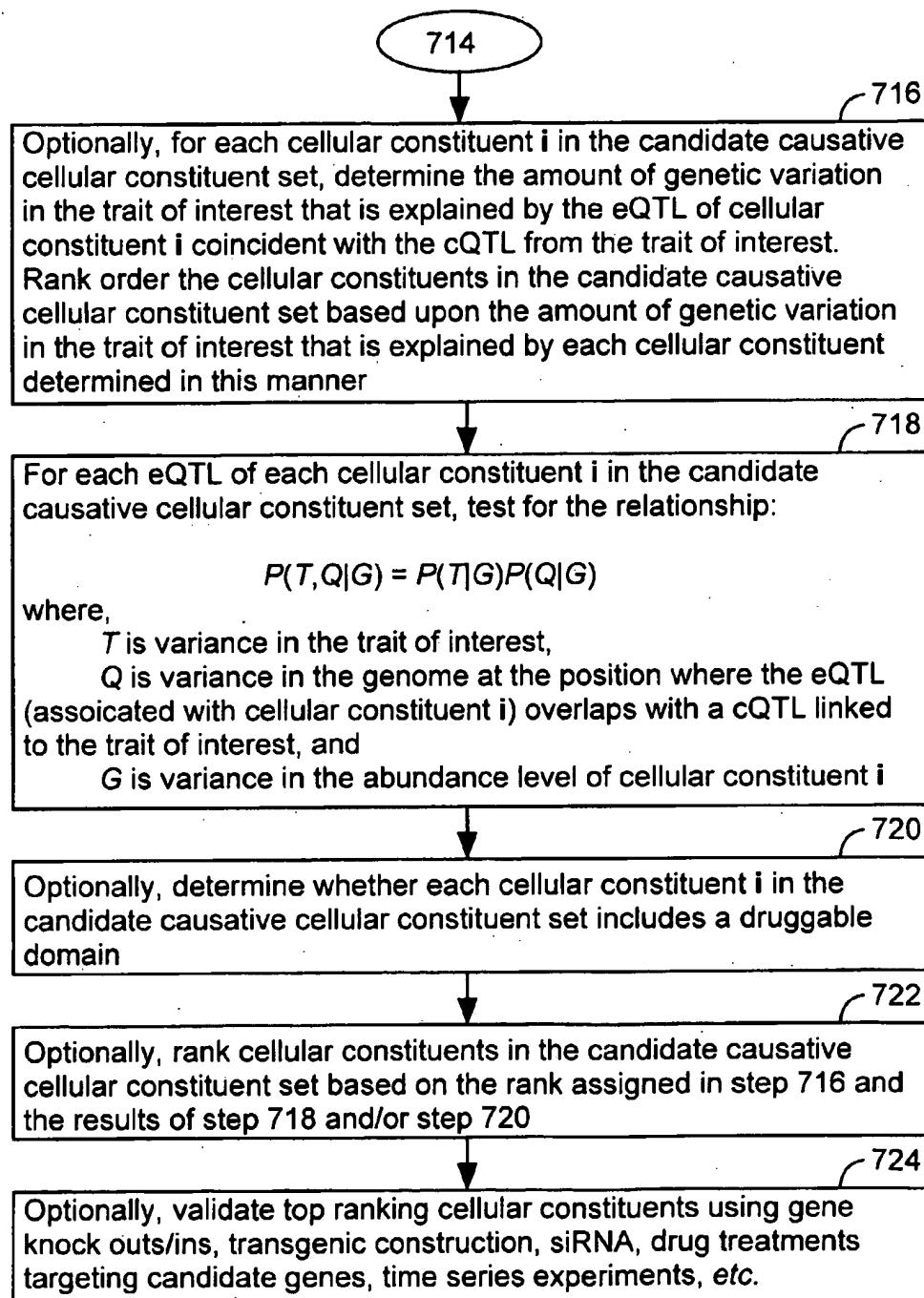
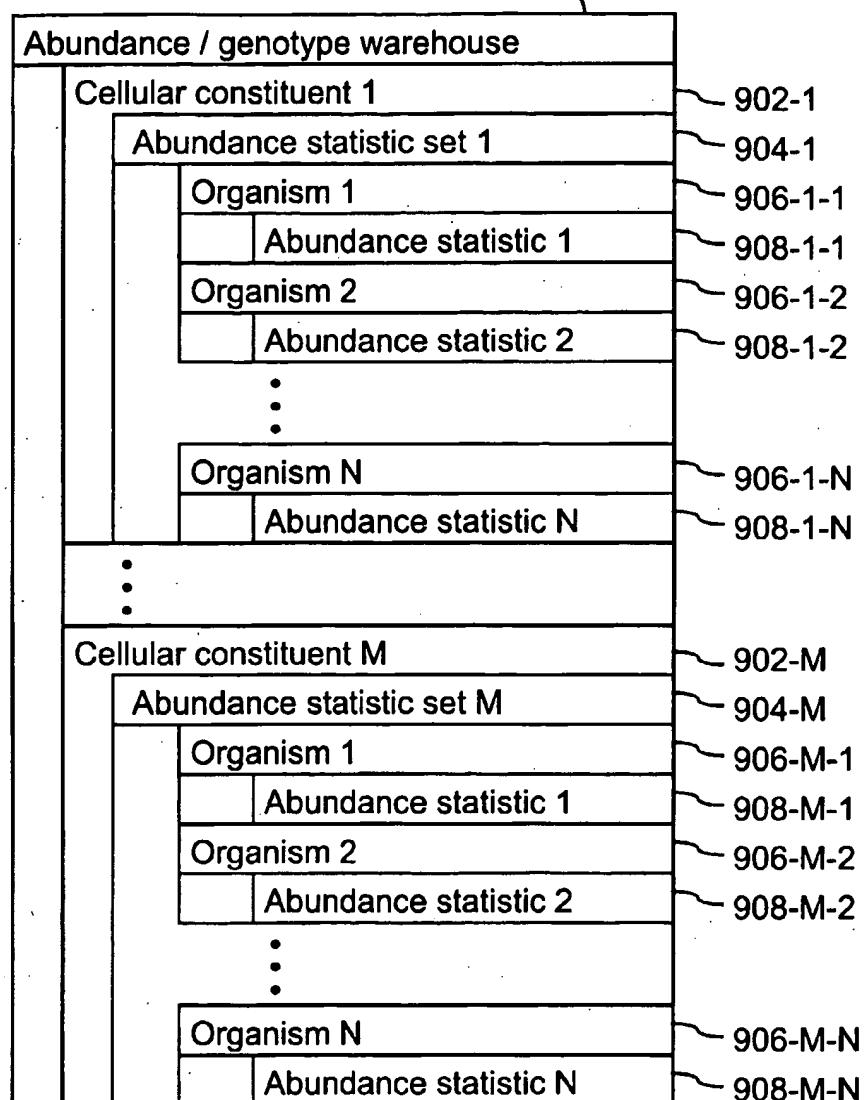


FIG. 7B

|   |         |
|---|---------|
| Phenotypic statistic set for clinical trait 1 |         |
| Phenotypic value for organism 1               | 74-1    |
| Phenotypic value for organism 2               | 804-1-1 |
| Phenotypic value for organism 3               | 804-1-2 |
| ⋮   | 804-1-3 |
| Phenotypic value for organism Q               | 804-1-Q |
| ⋮   |         |
| Phenotypic statistic set for clinical trait Z | 74-Z    |
| Phenotypic value for organism 1               | 804-Z-1 |
| Phenotypic value for organism 2               | 804-Z-2 |
| Phenotypic value for organism 3               | 804-Z-3 |
| ⋮   |         |
| Phenotypic value for organism Q               | 804-Z-Q |

**FIG. 8**

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**FIG. 9**

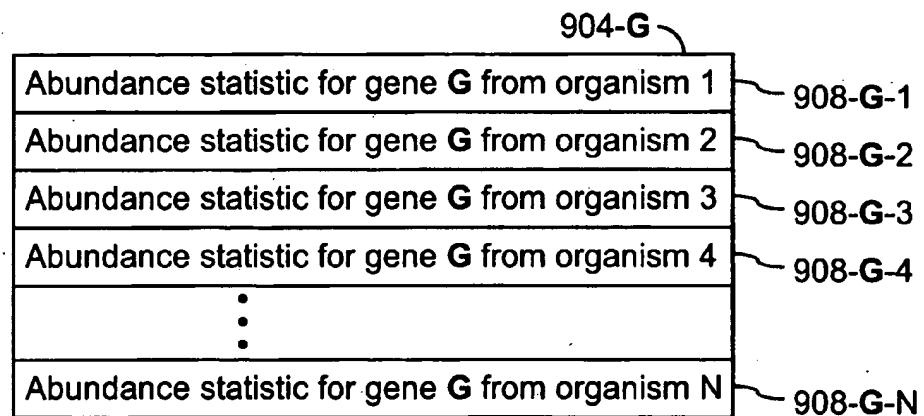


FIG. 10

78

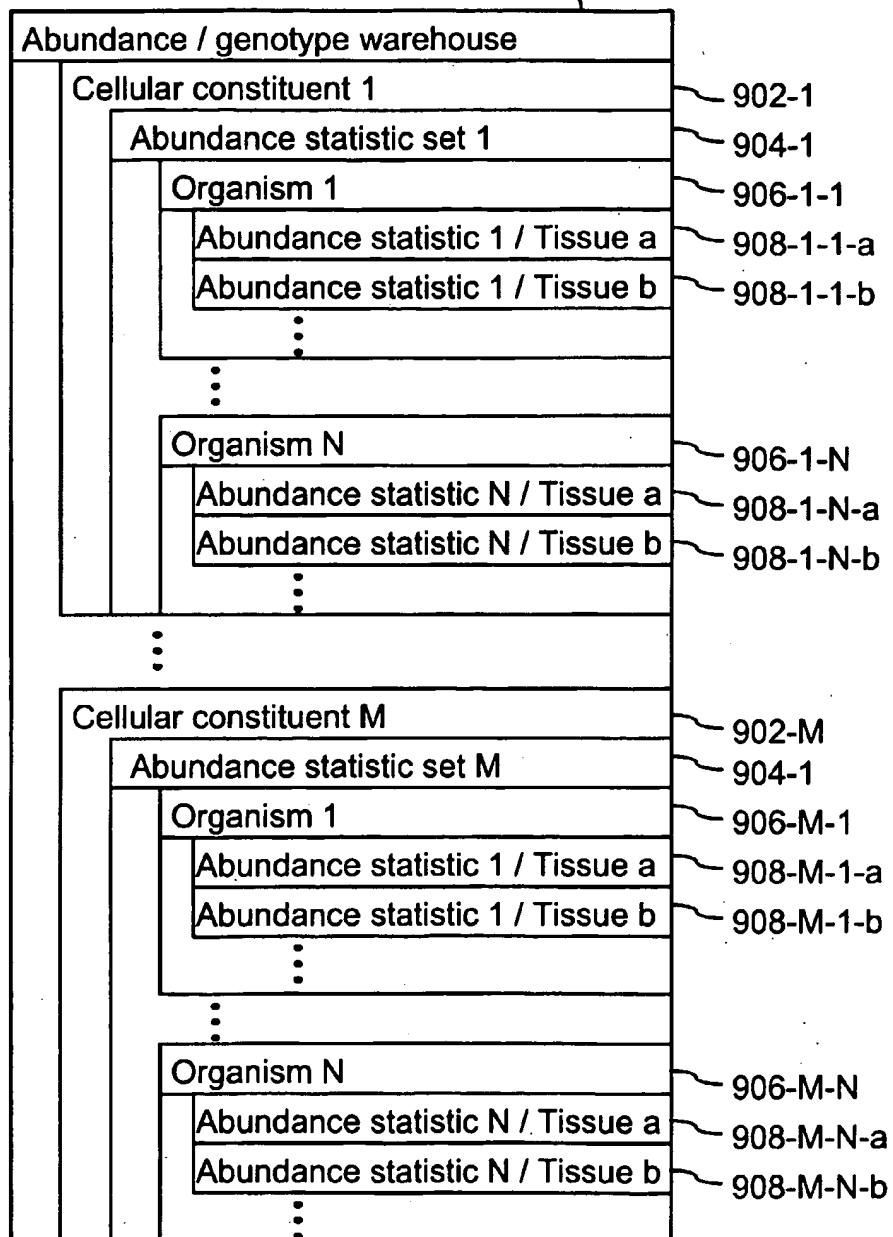
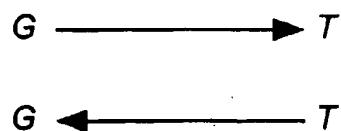
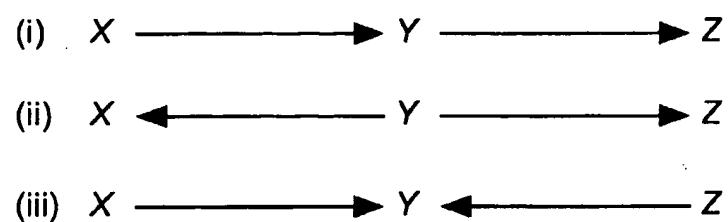
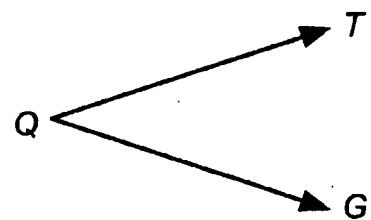


FIG. 11

| Abundance statistic set |                   |          |
|-------------------------|-------------------|----------|
| Position 1              |                   | 904-1    |
|                         | Statistical score | 1204-1-1 |
| Position 2              |                   | 1206-1-1 |
|                         | Statistical score | 1204-1-2 |
| ⋮                       |                   | 1206-1-2 |
| Position X              |                   | 1204-1-X |
|                         | Statistical score | 1206-1-X |
| ⋮                       |                   | ⋮        |
| Abundance statistic set |                   |          |
| Position 1              |                   | 904-M    |
|                         | Statistical score | 1204-M-1 |
| Position 2              |                   | 1206-M-1 |
|                         | Statistical score | 1204-M-2 |
| ⋮                       |                   | 1206-M-2 |
| Position X              |                   | 1204-M-X |
|                         | Statistical score | 1206-M-X |
| ⋮                       |                   | ⋮        |

**FIG. 12**

**FIG. 13A****FIG. 13B****FIG. 13C**

**FIG. 13D****FIG. 13E**

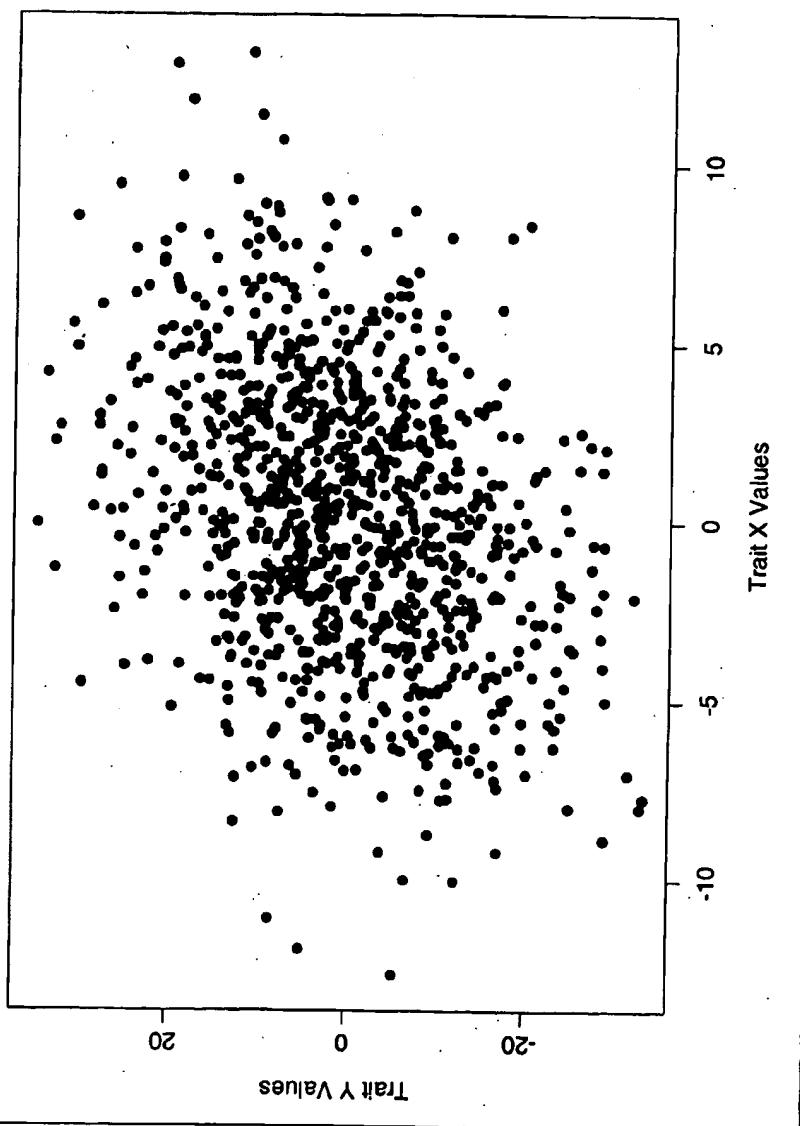
Scatter plot for traits X and Y ( $R^2 = 0.10$ )

Fig. 14

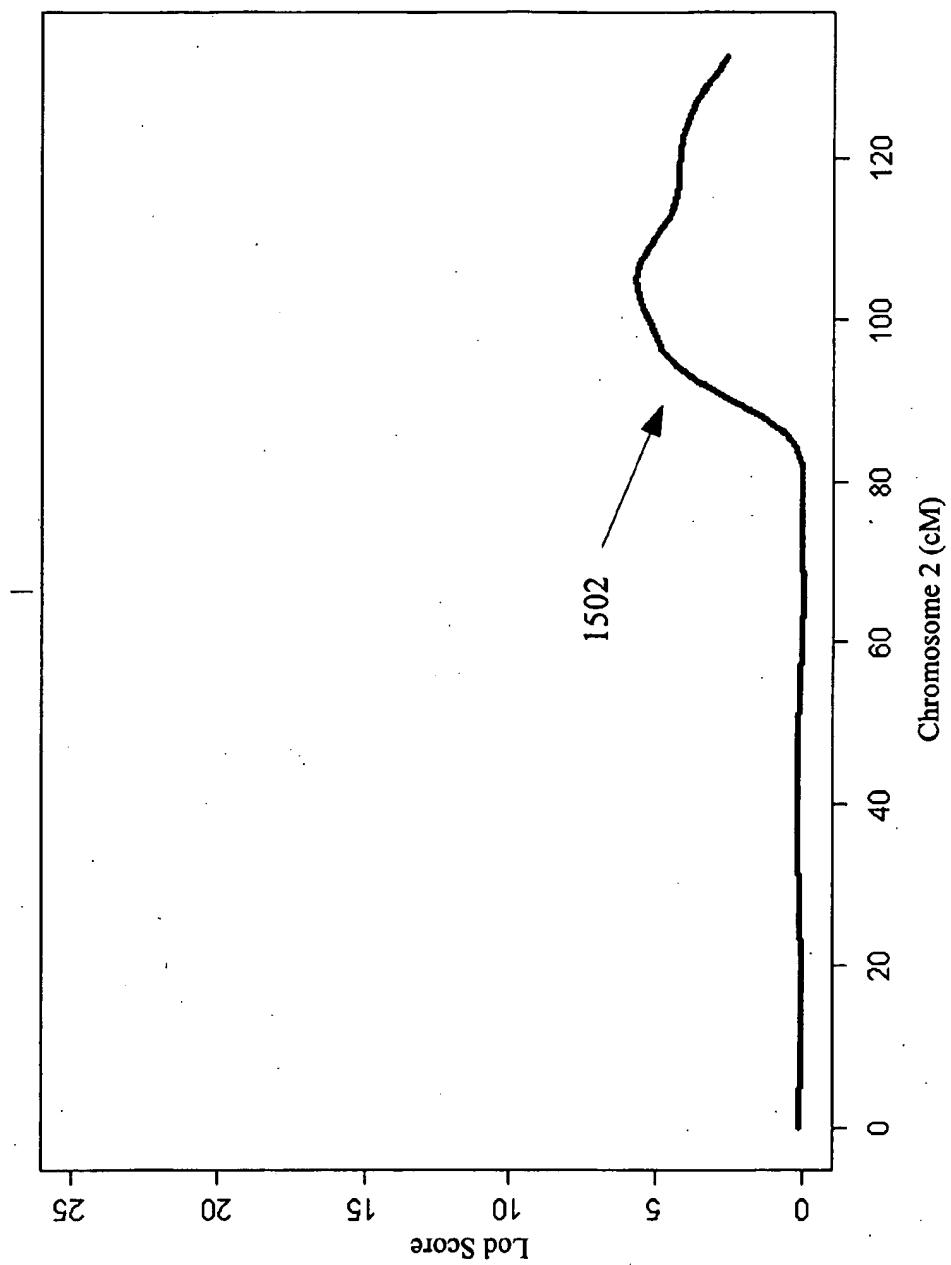
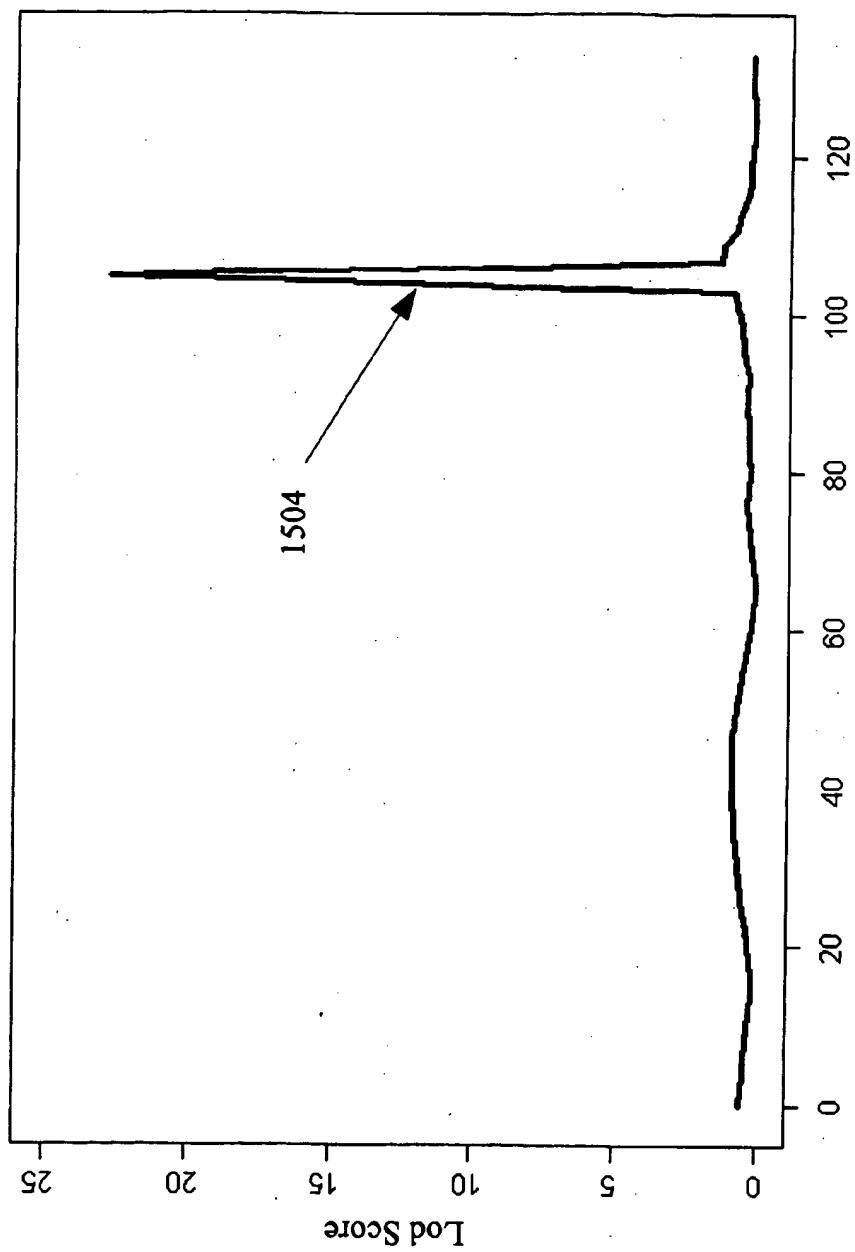


Fig. 15A



Chromosome 2 (cM)  
Fig. 15B

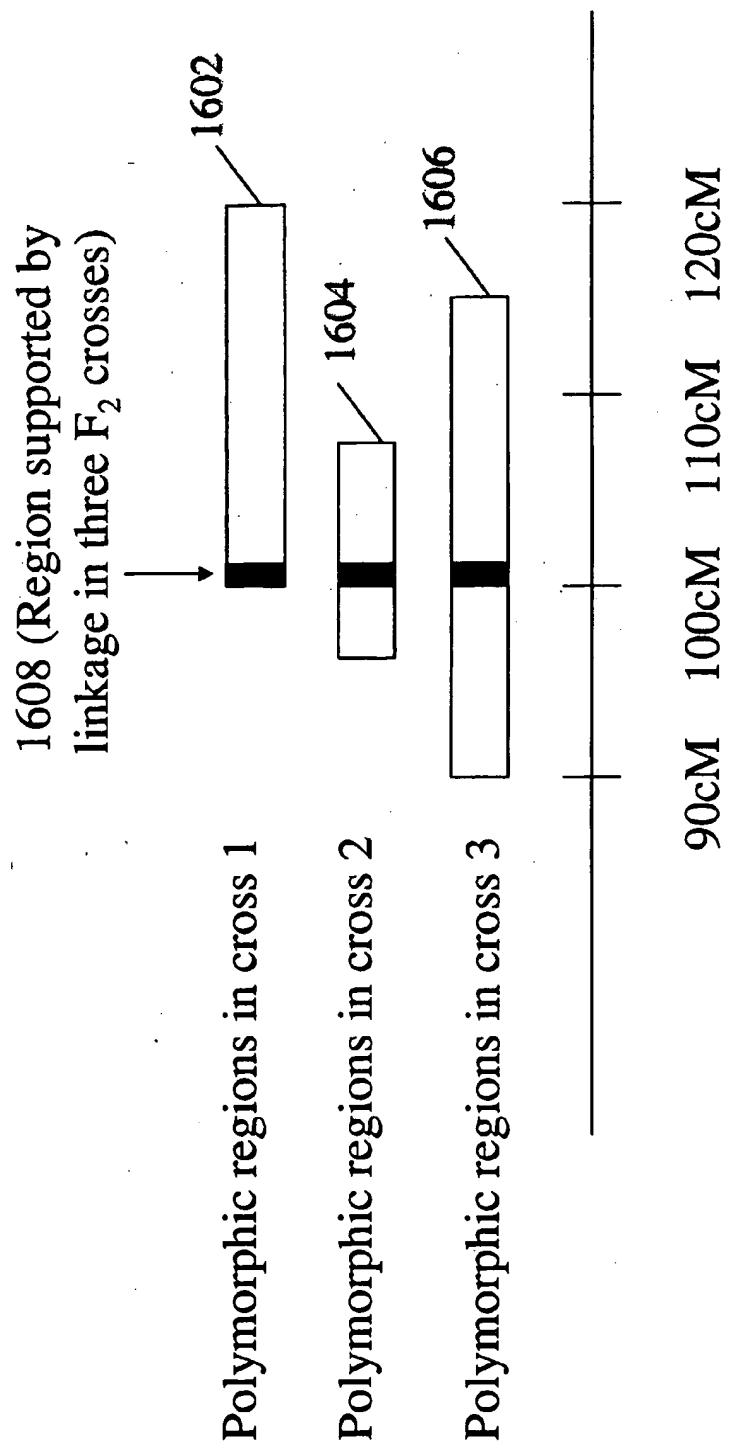


Fig. 16

|            |            |            |            |             |            |
|------------|------------|------------|------------|-------------|------------|
| 10         | 20         | 30         | 40         | 50          | 60         |
| MEPEAPRRRH | THQRGYLLTR | NPHLNKDLAF | TLEERQQLNI | HGLLPPSFNS  | QEIQVLRVVK |
| 70         | 80         | 90         | 100        | 110         | 120        |
| NFEHLNSDFD | RYLLLMDLQD | RNEKLFYRVL | TSDIEKFMPI | VYTPPTVGLAC | QQYSLVFRKP |
| 130        | 140        | 150        | 160        | 170         | 180        |
| RGLFITIHDR | GHIASVLA   | PEDVIKAI   | TDGERILGLG | DLGCNGMGIP  | VGKLALYTAC |
| 190        | 200        | 210        | 220        | 230         | 240        |
| GGMNPQECLP | VILDVGTE   | ELLKDPLYIG | LRQRRVRGSE | YDDFLDEFME  | AVSSKYGMNC |
| 250        | 260        | 270        | 280        | 290         | 300        |
| LIQFEDFANV | NAFRLLNKYR | NQYCTFNDDI | QGTASVAVAG | LAAALRITKN  | KLSDQTILFQ |
| 310        | 320        | 330        | 340        | 350         | 360        |
| GAGEAALGIA | HLIVMALEKE | GLPKEKA    | IWLVD      | VKG         | RASLTQE    |
| 370        | 380        | 390        | 400        | 410         | 420        |
| MKNLEAIVQE | IKPTALIGVA | AIGGAFSE   | QI         | PIIFALS     | NPT        |
| 430        | 440        | 450        | 460        | 470         | 480        |
| YKITKGRAIF | ASGSPFD    | LPNGQTLY   | PG         | VALGVVAC    | GL         |
| 490        | 500        | 510        | 520        | 530         | 540        |
| TAEVIAQQVS | DKHLEEGR   | Y          | LKIAEKIVKD | AYQEKTATVY  | PEPQNKEAFV |
| 550        | 560        | 570        |            |             |            |
| RSQMYSTDYD | QILPDCYSWP | EEVQKIQTKV | DQ         |             |            |

(SEQ ID NO: 1)

Fig. 17

|            |             |             |             |             |            |
|------------|-------------|-------------|-------------|-------------|------------|
| 10         | 20          | 30          | 40          | 50          | 60         |
| MEPRAPRRRH | THQRGYLLTR  | DPHLNKDLAF  | TLEERQQLNI  | HGLLPPCIIS  | QELQVLRIIK |
| 70         | 80          | 90          | 100         | 110         | 120        |
| NFERLNSDFD | RYLLLMDLQD  | RNEKLFYSVL  | MSDVEKFMP   | VYTPTVGLAC  | QQYSLAFRKP |
| 130        | 140         | 150         | 160         | 170         | 180        |
| RGLFISIHDK | GHIASVLA    | PEDVVKAIVV  | TDGERILGLG  | DLGCNGMGIP  | VGKLALYTAC |
| 190        | 200         | 210         | 220         | 230         | 240        |
| GGVNPQQCLP | ITLDVGTENE  | ELLKDPLYIG  | LRHRRVRGPE  | YDAFLDEFME  | AASSKYGMNC |
| 250        | 260         | 270         | 280         | 290         | 300        |
| LIQFEDFANR | NAFRLLNKYR  | NKYCTFNNDDI | QGTASVAVAG  | LLAALRITKN  | KLSDQTVLFQ |
| 310        | 320         | 330         | 340         | 350         | 360        |
| GAGEAALGIA | HLVVMAMEKE  | GLSKENARKK  | IWLVDISKGLI | VKGGRASLTEE | KEVFAHEHEE |
| 370        | 380         | 390         | 400         | 410         | 420        |
| MKNLEAIVQK | IKPTALIGVA  | AIGGAFTEQI  | LKDMAAFNER  | PIIFALSSPT  | SKAECSADEC |
| 430        | 440         | 450         | 460         | 470         | 480        |
| YKVTKGRAIF | ASGSPFPDPVT | LPDGRTLFPG  | QGNNSYVFPG  | VALGVVACGL  | RHIDDKVFLT |
| 490        | 500         | 510         | 520         | 530         | 540        |
| TREVISQQVS | DKHLQEGRLY  | PPLNTIRGV   | LKIAVKIVQD  | AYKEKMATVY  | PEPQNKEEFV |
| 550        | 560         | 570         |             |             |            |
| SSQMYSTNYD | QILPDCYPWP  | AEVQKIQTKV  | NQ          |             |            |

(SEQ ID NO: 2)

Fig. 18

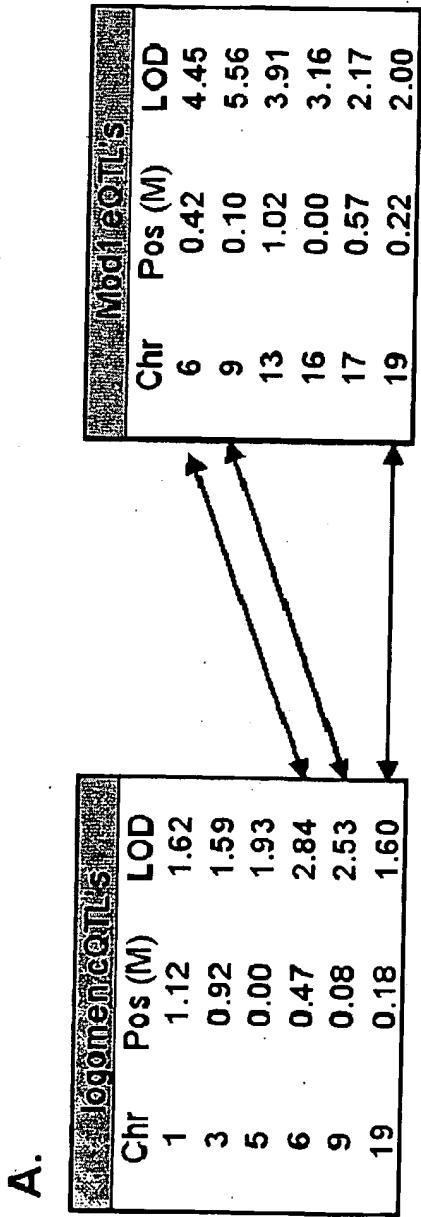


Fig. 19A

**B.**

| No. of Overlaps With Model |   |
|----------------------------|---|
| logomen                    | 3 |
| epipa                      | 2 |
| flpsum                     | 2 |
| lep                        | 2 |
| logflpsum                  | 2 |
| logsubc                    | 2 |
| sqrtipipa                  | 2 |
| sqrtlep                    | 2 |

| No. of Overlaps With Model |   |
|----------------------------|---|
| sqrtretrog                 | 2 |
| fatbw                      | 1 |
| livebw                     | 1 |
| omen                       | 1 |
| subc                       | 1 |
| irs                        | 0 |
| retrog                     | 0 |

Fig. 19B

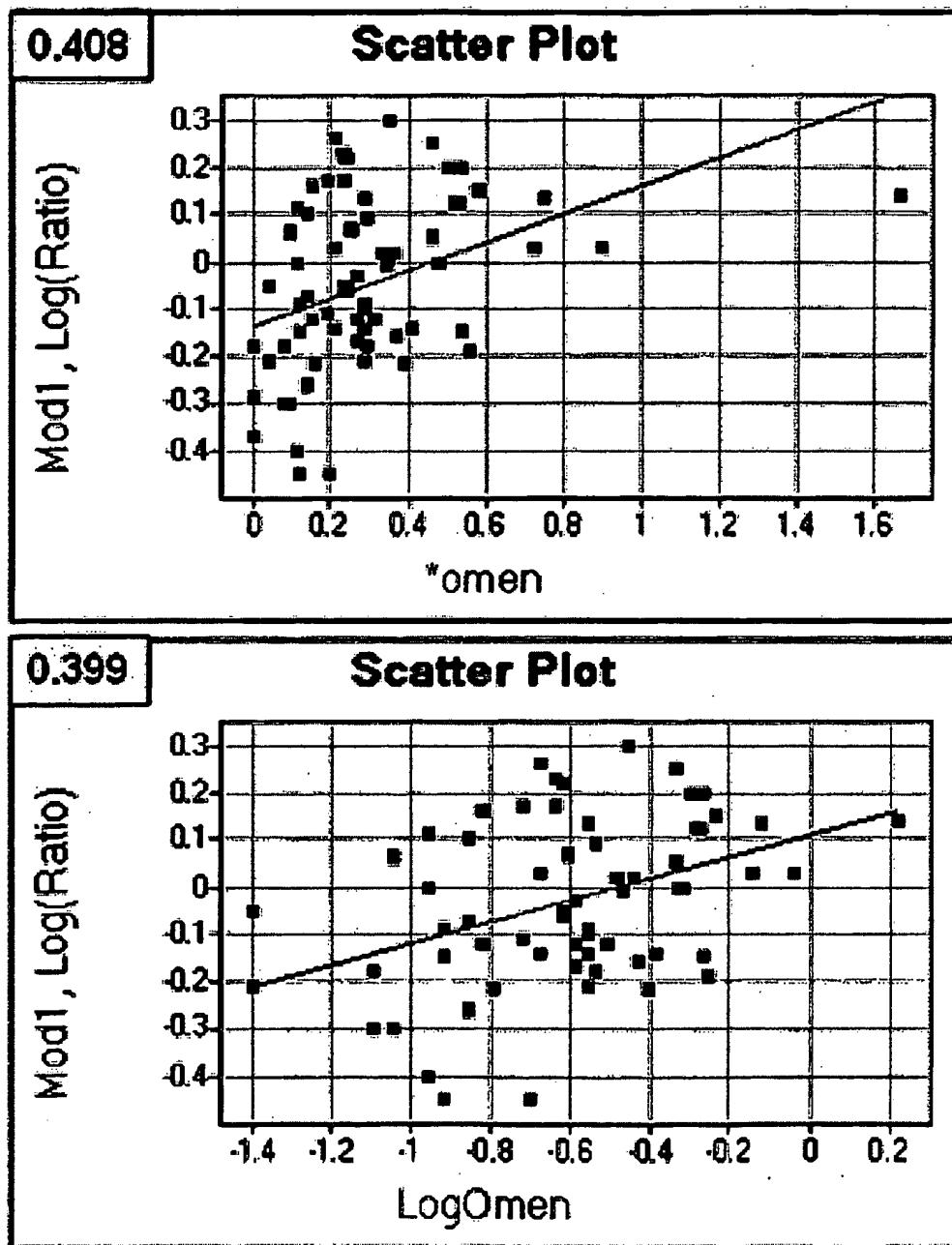


Fig. 20

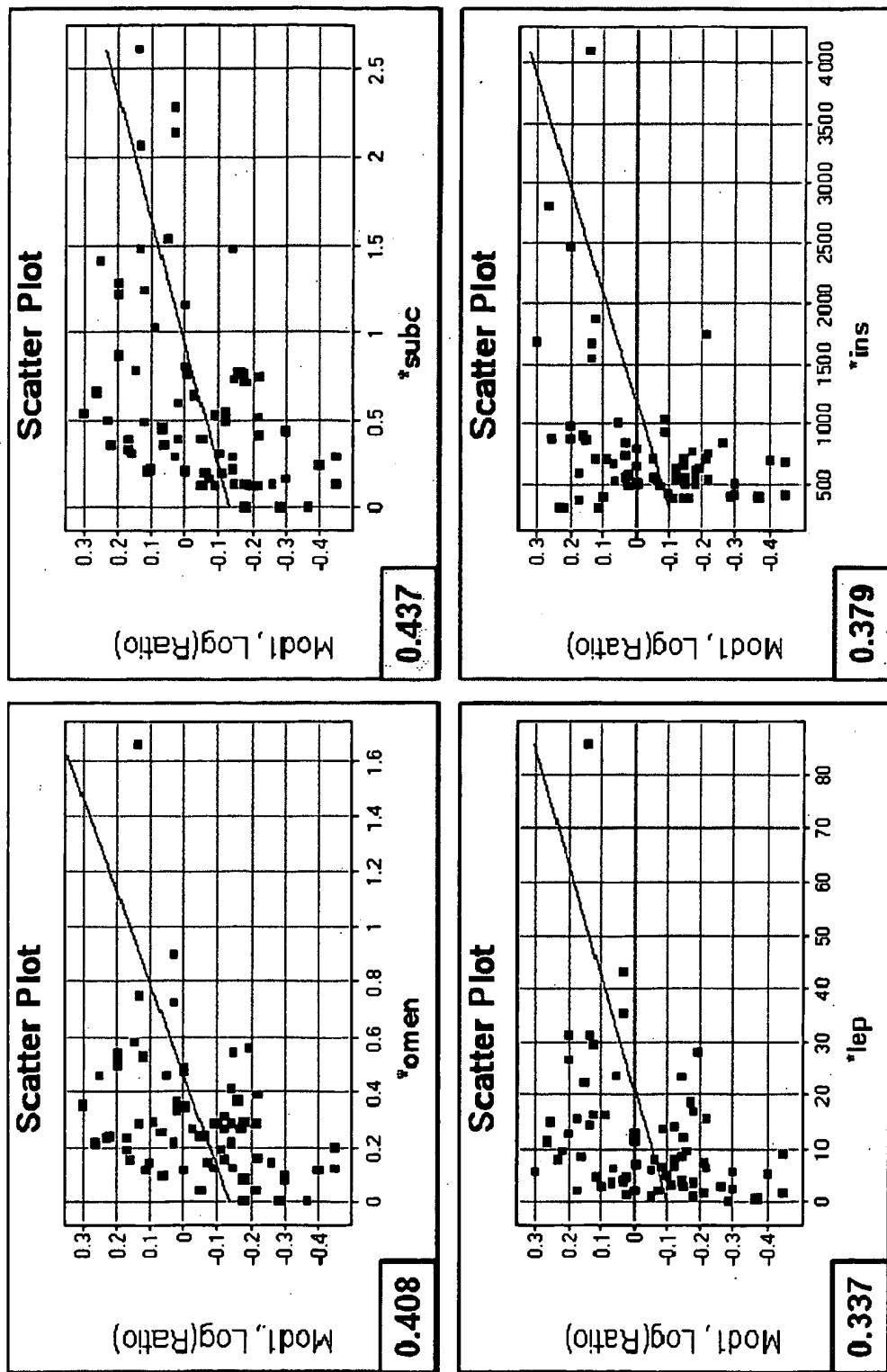


Figure 21

|          | *livebwt | *retrog | *epipa | *omen | *subc | *fipsum | *fatbw | *lep | Mod1 |
|----------|----------|---------|--------|-------|-------|---------|--------|------|------|
| *livebwt | 1        | 0.56    | 0.65   | 0.64  | 0.62  | 0.67    | 0.43   | 0.67 | 0.23 |
| *retrog  |          | 1       | 0.77   | 0.78  | 0.75  | 0.82    | 0.78   | 0.76 | 0.49 |
| *epipa   |          |         | 1      | 0.89  | 0.86  | 0.99    | 0.91   | 0.94 | 0.36 |
| *omen    |          |         |        | 1     | 0.84  | 0.92    | 0.82   | 0.92 | 0.41 |
| *subc    |          |         |        |       | 1     | 0.92    | 0.87   | 0.85 | 0.44 |
| *fipsum  |          |         |        |       |       | 1       | 0.92   | 0.95 | 0.41 |
| *fatbw   |          |         |        |       |       |         | 1      | 0.82 | 0.45 |
| *lep     |          |         |        |       |       |         |        | 1    | 0.34 |
| Mod1     |          |         |        |       |       |         |        |      | 1    |

Figure 22

1 IKEKGKPLXL NPRTNKGXAF TLQERQXLGL QGLLPPKIEQ QDIQALRFHR  
51 NLKKXTSPLK KYIYIXGIQE RNEKLFYRIL QDDIESLXPI VYTPTVGLAC  
101 SQYGHIFRRP KGLFISISDR GHVRSIVDNW PENHVKAVVV TDGERILGLG  
151 DLGVYGXGIP VGKLCLYTAC AGIRPDRCLP VCIDVGTDNI ALLKDPFYXG  
201 LYQKRDRTQQ YDDLIDEFXK AITDRYGRNT LIQFEDFGNH NAFRFLRKYR  
251 EKYCTFNDDI QGTAVALAG LLAAQKVISK PISEHKILFL GAGEAALGIA  
301 NLIVXSXVEN GLSEQEAQKK IWXFDKYGLL VKGRKAKIDS YQEPFTHSAP  
351 ESIPDTFEDA VNILKPSTII GVAGAGRLFT PDVIRAXASI NERPVIFALS  
401 NPTAQAECTA EEAYTLTEGR CLFASGSPFG PVKLTDGRVF TPGQGNVYI  
451 FPGVALAVIL CNTRHISDSV FLEAAKALTS QLTDEELAQG RLYPPLANIQ  
501 EVSINIAIKV TEYLYANKXA FRYPEPEDKA KYVKERTWRS EYDSLLPDVY  
551 EWPESSASSPP VITE

(SEQ ID NO: 3)

Fig. 23

|             |            |            |            |             |            |
|-------------|------------|------------|------------|-------------|------------|
| 10          | 20         | 30         | 40         | 50          | 60         |
| MLSRLRVVST  | TCTLACRHLH | IKEKGKPLML | NPRTNKGMAT | TLQERQMLGL  | QGLLPPKIET |
| 70          | 80         | 90         | 100        | 110         | 120        |
| QDIQALRFHR  | NLKMKMTSPL | KYIYIMGIQE | RNEKLFYRIL | QDDIESLMPI  | VYTPTVGLAC |
| 130         | 140        | 150        | 160        | 170         | 180        |
| SQYGHIFRRP  | KGLFISISDR | GHVRSIVDNW | PENHVKAVVV | TDGERILGLG  | DLGVYGMGIP |
| 190         | 200        | 210        | 220        | 230         | 240        |
| VGKLCLYTAC  | AGIRPDRCLP | VCIDVGTDNI | ALLKDPFYMG | LYQKRDRTQQ  | YDDLIDEFMK |
| 250         | 260        | 270        | 280        | 290         | 300        |
| AITDRYGRNT  | LIQFEDFGNH | NAFRFLRKYR | EKYCTFNDDI | QGTAVALAG   | LLAAQKVISK |
| 310         | 320        | 330        | 340        | 350         | 360        |
| PISEHKILFL  | GAGEAALGIA | NLIVMSMVEN | GLSEQEAQKK | IWMFDKYGLL  | VKGRAKIDS  |
| 370         | 380        | 390        | 400        | 410         | 420        |
| YQEPFTHSAP  | ESIPDFTEDA | VNILKPSTII | GVAGAGRLFT | PDVIRAMASI  | NERPVIFALS |
| 430         | 440        | 450        | 460        | 470         | 480        |
| NPTAQAEACTA | EEAYTLTEGR | CLFASGSPFG | PVKLTDGRVF | TPGQGNVYI   | FPGVALAVIL |
| 490         | 500        | 510        | 520        | 530         | 540        |
| CNTRHISDSV  | FLEAAKALTS | QLTDEELAQG | RLYPPLANIQ | EV SINIAIKV | TEYLYANKMA |
| 550         | 560        | 570        | 580        |             |            |
| FRYPEPEDKA  | KYVKERTWRS | EYDSLIPDVY | EWPESASSPP | VITE        |            |

(SEQ ID NO: 4)

Fig. 24

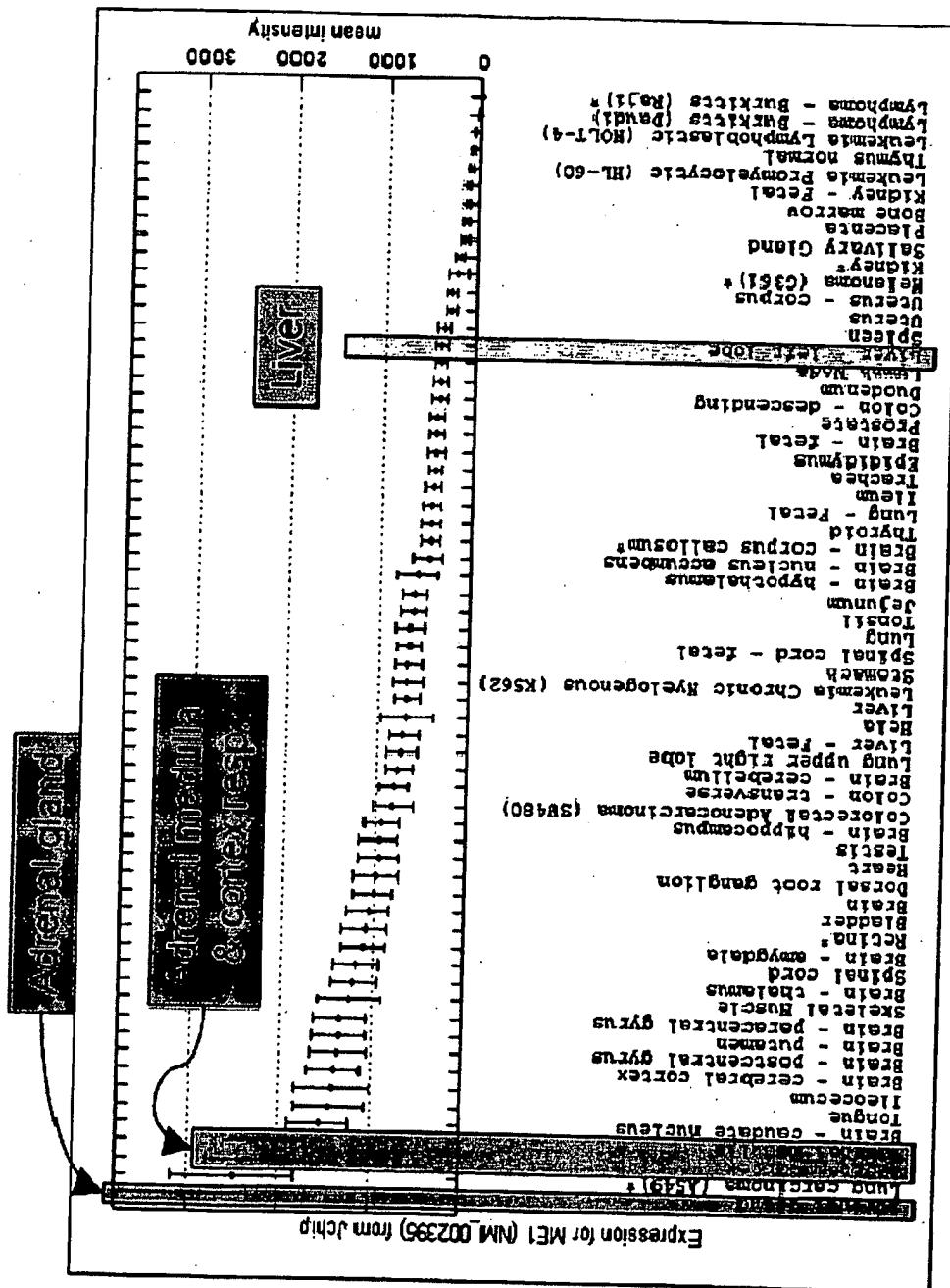


Fig. 25

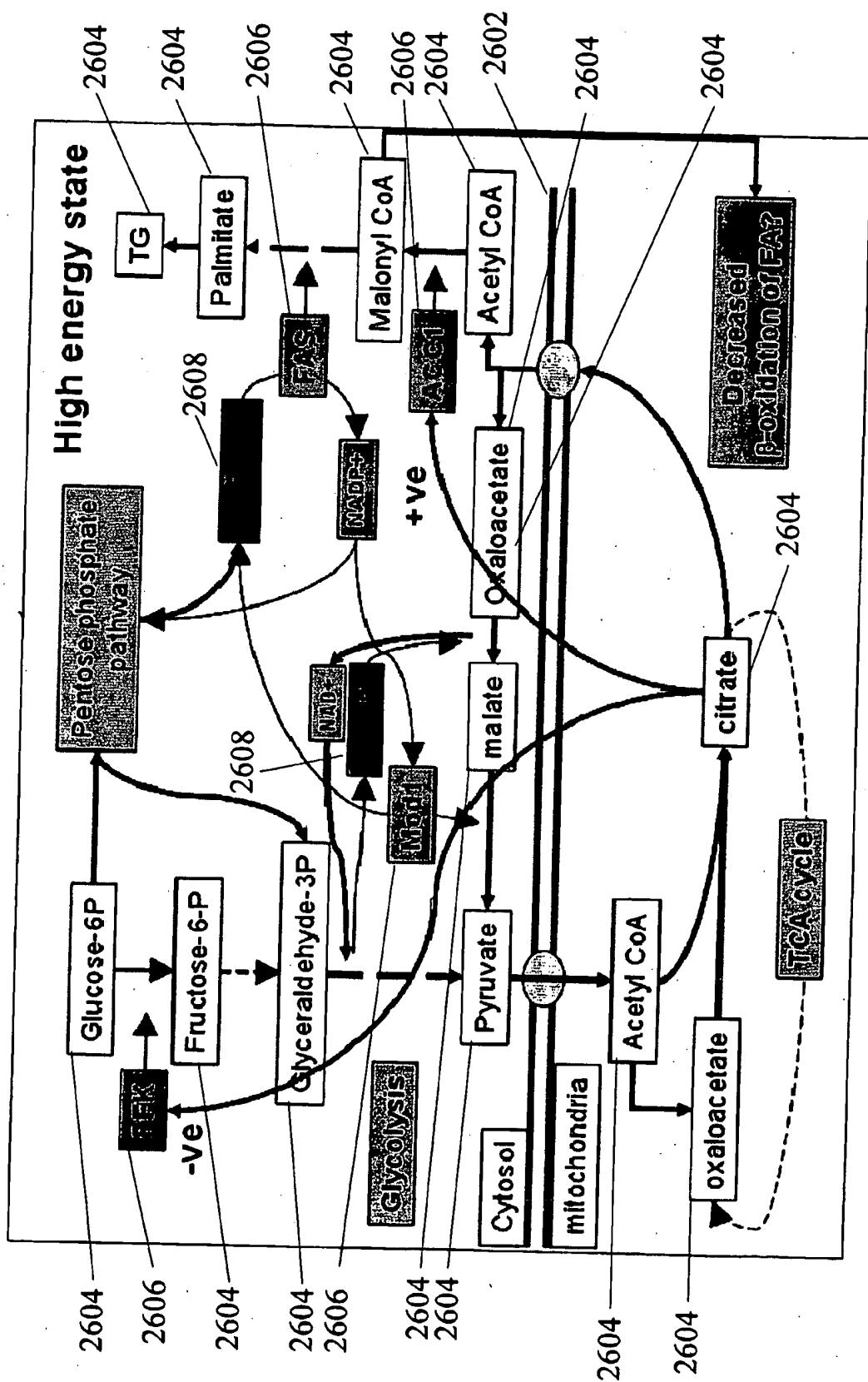


Fig. 26

1 atggcctta cccttgaaga aaggctgcag ctggaatcc acggccta at cccgcctgc  
61 tttctgagcc aggacgtcca gtcctccga atcatgagat attacgagcg gcagcagagt  
121 gacctggaca agtacatcat tctcatgaca ctccaagacc gtaacgagaa gctcttctac  
181 cgagtgctga cttcggacgt ggagaagttc atgccaatcg tgtacacgcc taccgtgggg  
241 ctagcctgtc agcactatgg cctgacttcc cgccaggcccc gtggactgtt catcaccatt  
301 catgacaaag gtcatcttgc aacaatgctg aattcttggc cagaagacaa tattaaggcc  
361 gtgggtgtga ctgatggggc ggcacatctg ggcctggag acctggctg ctacggcatg  
421 ggcacatccctg tggcaagct ggcctgtac acggcatgcg gaggggtgaa cccgcagcag  
481 tgcctccctg tgctgctgga cgtcggcacc aacaatgagg agctgctcag agaccctctg  
541 tacatcgcc taaaacacca ggcgtgcac gggaggcat acgatgactt gctggatgag  
601 ttcatgcagg ctgtgacaga caagtttggc ataaattgcc tcacccaatt tgaagacttc  
661 gccaatgcca atgccttccg cctgctcaac aaataccgta acaagtactg catgttaat  
721 gatgacatcc aagatgactt ctccagaggc ccaaagaggt cacaactttt cttcaagtga

(SEQ ID NO: 5)

Fig. 27

1 atgttgcg ggttaagcgt agttccacc acttgtactt tggcatgtct acatttacac  
61 ataaaaagaaa aaggcaagcc acttatgctg aatccaagaa caaacaaggg aatggcattt  
121 actttacaag aacgacagat gcttggctt caagggctt tacctccccca aatacagaca  
181 taagatattc aaggcttacg attccataga aacttgaaaaaa aatgactag cccttcggaa  
241 aactatatct acataatggg aatacaagaa agaaatgata aattgtttta tagaatactg  
301 caagatgaca cggagagttt aatgccaatt gcatatacac cgacgggtgg tcttgtctgc  
361 tcccagtgtg gacaccttctt tagaagacct aagggattat ttatccat ctcagacaga  
421 ggtcatgtt gatcaattgtt ggataagtgg ccagaaaaatc atgttaaggc tgtttagtg  
481 actgatggag agagaattct gggtcatgga gatctgggtg tctatggat gggattcca  
541 gtagaaaaaa ttgtttgtt tacagttgtt ccaggaatat ggcctgatag atgccttctg  
601 gtgtgtattt atgtgggagc tgataatatc gcactcttaa aaggcacatt ttacatggc  
661 ttgtaccaga aacgagatcg cacacaacag tctgatgatc caattgtatgaa gtttatgaaa  
721 gctattactg acagatatgg ctgaaacaca ctcccttcagt ttgaaggttt tggacatcat  
781 aatgcattca gattctttagg aaaataccaa taaaaatgtt gcactttcaa tgatgatatt  
841 caagggacag ctgcagtagc tctaatacggtt ctcttgcaa cacaaaaagt tactagtaaa  
901 ccaatctccg aacacaaaaat cttattccctt ggagcaggag agattactt tagaattgca  
961 aatctttag tattgtctat ggttagaaaaat ggccctgtcag aagaagaggc aaaaaagaaa  
1021 atctggatgt ttgacaagta tggtttatta gtttaggggc agaaagcaaa aatagattgt  
1081 ttcaggaaac catttactta cccagtcctt cttttttttt gatctttttt tttttttttt  
1141 gtgaatataa tgaagacttc aactacaatt ggagttgcag gtgtggccg tcttttact  
1201 cctgatgtaa tcagagccat tggctgtatc aatgaaaggc ctgtatattt tgatattttt  
1261 aatcttacag cacaggccgg gtcacggctg gagaagcata tacacttaca  
1321 gagggcaaat gtttgggttgc cagtggcagt ccattttggc cagtgaaact cacagatggg  
1381 cgaatcttta caccagatcg aggaaacaaat gtatataattt ttccaggtgt gacttttagct  
1441 gttattctct tgaacacccca gcaaatttagt gacaatgttt tccttagaagc tgcaaaggca  
1501 ttgacaagcc acgtgacggg tgacgcgtt gcccgggggaa gactttactt accacttgct  
1561 aatattcaga aagtttctat taacattgtt ataaaaatgtt cagaataacct gtatgttaat  
1621 aaaaatggctt tctcaataacc cagaacacctga

(SEQ ID NO: 6)

Fig. 28

1 ccggccgac agctgcagtc agcaccgtca ccccagcagc atccgcccgc tgacccgc  
61 gtgcggcccg cccccggctcg acccccgcgc cgaacccggc gccagccatg gagcccgaag  
121 cccccccgtcg cggccacacc catcagcgcg gctacctgtc gacacggAAC cctcacctca  
181 acaaggactt ggccttacc ctggaaagaga gacagcaatt gaacattcat ggattgttgc  
241 caccccttca acacagtca gagatccagg ttcttagagt agtaaaaaat ttcgagcatc  
301 tgaactctga ctttgacagg tatcttctct taatggatct ccaagataga aataaaaac  
361 tctttatag agtgctgaca tctgacattt agaaaattcat gcctattgtt tataactccca  
421 ctgtgggtct ggcttgccaa caatatagtt tggtgttgc gaagccaaaga ggctcttta  
481 ttactatcca cgatcgaggg catattgtt cagttctcaa tgcatggcca gaagatgtca  
541 tcaaggccat tgggtgact gatggagac gtattcttgg ctggagac ctggctgt  
601 atggaatggg catccctgtg ggttaaattgg ctctatatac agcttgcggg gggatgaatc  
661 ctcagaatg tctgcctgtc attctggatg tggttgcggaa aatgaggag ttacttaaag  
721 atccactcta cattggacta cggcagagaa gagtaagagg ttctgaatat gatgatttt  
781 tggacgaatt catggaggca gtttcttca agatggcat gaattgcctt attcagtttgc  
841 aagatttgc caatgtgaat gcatttcgtc tcctgaacaa gtatcgaaac cagtattgca  
901 cattcaatga tgatattcaa ggaacacgt ctgttgcagt tgcatggctc ctgcagctc  
961 ttcaaaaatcaac caagaacaaa ctgtctgatc aaacaataact attccaagga gctggagagg  
1021 ctgccttagg gattgcacac ctgattgtga tggccttggaaa aaaaaggt ttaccaaaag  
1081 agaaagccat caaaaagata tggctgggtt attcaaaagg attaataatgtt aaggacgtg  
1141 cttcccttaac acaagagaaa gagaagtttt cccatgaaca tgaagaaatg aagaacctag  
1201 aagccattgt tcaagaaaatc aaaccaactg ccctcatagg agttgctgca attgggtgg  
1261 catttcaga acaaatttctc aaagatatgg ctgccttcaa tgaacggctt attattttgc  
1321 cttttagtaa tccaaacttagc aaagcagaat gtctgcaga gcagtgcataaaaataacca  
1381 agggacgtgc aatttttgcg agtggcgtc cttttgcattc agtcaactt ccaaataatgg  
1441 agacccata tcctggccaa ggcacaaatt cctacgtgtt ccctggagtt gctttgg  
1501 ttgtggcgtg tggattgagg cagatcacag ataataatttt cctcaactt gctgagg  
1561 tagtcagca agtgcagat aaacacttgg aagagggtcg gctttatcctt ctttgcata  
1621 ccatttagaga tttttctctg aaaaattgcg aaaaaggatgt gaaagatgc taccaaaa  
1681 agacagccac agtttacccat gaaacggcaaa acaaagaagc atttgcgc tcccaatgt  
1741 atagactga ttatgaccag atttacccat attgttattt ttggcctgaa gaggtgcaga  
1801 aaatacagac caaagttgac cagtaggata atagcaaaaactt tttcaactc tattaatgag  
1861 gtctttaaac ctttcataat tttaaagggt tggatcttt tataatgatt cataagacac  
1921 ttagattaag attttactt aacagtcataa aaattgtatag aagaataatcg atataaattg  
1981 ggataaaacat cacatgagac aattttgcctt cactttgcct tctggatttattatg  
2041 gtctgaatta ttctgcctac gttctttta aaagctgtt tacgtactac ggagaaactc  
2101 atcattttta tacaggacac taatggaaag accaaaatta ctaataaattt gaaataacca  
2161 acattaaaac tcataattat tttgttgcattt aatctacttt tc

(SEQ ID NO: 7)

Fig. 29

CCTGAAAACACTTATAACGGGGTAGGGCAATTATACATAGCAAACCCGTAAACATTT  
AAACTCTAATTAAACATTTAATTCTTCAGAATTAAATACACACAGATGCTATCATGGGG  
GGGGGAGGCGTGCAGCCCCATCGGACCGGGTTTGACAACTTACTTACAACATTATTA  
CATCCTTTATTACTGGTCCAGGCCCGGAGCATGGAAAGATATACAGCGTGGAGTAAA  
CACATTCATCCTGGGTGAGGAGTTCTGGCAGGAGACACTGCTTTCAACATTAAAAATGT  
ATAAGGTGTTAGCAAAAGTTACAGAAAACGGACCAAATGAGCAAGTTATTTGTTAGA  
AAATTCCACTTCGTGGGTTCGCTGATGTGCTCGGGTTGCAAGGGAATGCTTCCGG

(SEQ ID NO: 8)

Fig. 30

1 gttgcagagc agtactgcgg ggaacaagaa actgcagcgg gcgctagagg ggcggacctg  
 61 aggtcgcgga ttccgaagcc ccggaggcag attccgagtg cagtggtag gaggtgtcc  
 121 tccgggcctc gccgaccatc ctgcggacgg actgggcgtg gccggaggaa ctgtcccga  
 181 gctgtggggc ttccatggc gccttggaa gaggcaggcagg agaaggcggg gctccccc  
 241 acgttccggc cgaagtggct gcagagctga aggggtgggg cctcgggta gcccggtag  
 301 tggatctgt cctctctct cagccctgga ccatacgccag cacacactga ggcaggaatg  
 361 gccccgagac ctccgacggc caagccccag gagtcggtga cattcaaaga tggctgt  
 421 aacttcaccc aggaagaatg gcaccacgtg gcccctgccc agaggagctt atacagggat  
 481 gtcatgctgg agaactacaa ccacctggc tcgctcgggt atcaagtctc caagccagag  
 541 gtgatcttca aattggagca aggagaagag ccatacgat cagagaaaga aatccaaaga  
 601 ccttctgtc cagactggaa gaccaggcct gagtcctcac ggagtctca gcagggcgt  
 661 tctgaagtat tccctcagaac aaatgttta tcacacacca caataggtga tatctggaat  
 721 gtcgttatcc agggggcatca ggaaagtggg agaagacatc tggggccaga ggcacatcttcc  
 781 cagaagaaaa taaccactt agagaaaaaa attgagcaaa acaaaggtag tgaagactct  
 841 agtttgagca cagacttggc tccacaactg gacatttctt caagtataag gcccaggtag  
 901 tgtaaaacat ttggaaataa ttggaaacac aattcagaac tagttactca gagaatata  
 961 cttgttaaaa agaaggccta taagtgtat aaatgttagga aatcatttat tcatacatca  
 1021 tcacttaata aacacgagaa gattcataaa ggcgatccctt actccaatgg tacagacca  
 1081 ggagctcagt ctggaaaggaa acaccatgag tggcggact gtggaaaac cttccctctgg  
 1141 agaacacacgc tcacggagca ccagagaatt cacactgggg aaaaaccctt tgagtgtat  
 1201 gtgtgtggaa aggccttcag gcacagctcg tccttggc agcatggaaa cgacacataca  
 1261 ggagagaagc cctatcagt tagcctctgt gggaaaggcct tccagcgcag ttcatctt  
 1321 gttcaacacc agagaatcca cacggagag aagccctatc gtcataatc ctgtgggagg  
 1381 tcattcaggc acagcacgtc cctcagcga catgaggtag cccacagtgg ggagaaaccc  
 1441 ttccagtgtc aggaatgtgg gaaggcctt agcagggtgtt cttcccttgc ccagcatgag  
 1501 aggacccata caggagagaa gccttcgag tgccatgtt gtggggggc atttggtag  
 1561 agcccatccc ttataaaaca tatgaggatt cataaaagaa gcaaacctt ccaaagtaac  
 1621 aacttcagcc tggctttgt gcctaactt cctcttctc aaggtgaagg cctgcttact  
 1681 gaagtaaaat cgttccatgg taatgactgt gggaaagact tcggtcacat tacagacttc  
 1741 tctgagcacc agaggctcca cgctgggg aatttcctacg gtcataaca gacccctt  
 1801 ggtcagcagt ccctgtctca tccccggag aaaccctatc agtgcacatc atgtggaaa  
 1861 gctttaaaaa ggagtacaag tttatagag catcatcgaa ttcacactgg agagaaaccc  
 1921 tatgaatgtt atgagtgtgg ggaaggcctt agtgcactt cgtcactcac gcaacacgag  
 1981 aggacccaca ctggcgagaa accctatgag tgcatgtact gcgggaaagc cttcagtc  
 2041 agctcatccc tgattcagca cgaaaggacg cataccggag agaaacccta tgagtgtat  
 2101 gagttggc gggccttag aaagaagacc aatttgcacg accatcagag aactcacact  
 2161 ggagagaaac cctatgtttt caaggagtgt gggagaaact tcagccggag ctccggcc  
 2221 actaaacacc accgagttca cgccggaaat aaactgcagg aaagctaaac aatggatgg  
 2281 ggaggaggca cggccgaaca tctgttcca acccagtgtc agaggattct gaaagtctga  
 2341 gaatgttattt atgtgtttgg acactgtgt tagagaaaaac tgccactaga agaaaaaaat  
 2401 tttaaattaa agccatttctt tcatacctt ttacaggctt cttgttagaac tacgtacggc  
 2461 atatgttagtc gtttggaaat gatgtgacct tactaaagct tttgaatata tgggtgcaga  
 2521 gtcaccaagt tttaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa  
 2581 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa

(SEQ ID NO: 9)

Fig. 31

1 maprppptakp qesvtfkdva vnftqeewhh vpgaqrslyr dvmlenynhl vslgyqvskp  
 61 evifklegge epwisekei q rpfcpdwktr pessrspqgg vsevflrtvn lshttigdiw  
 121 nvaiqghqes grrhlgpeas sqkkittlek kieqnkvged sslstdlvpq ldiisssirps  
 181 dcktfgnnle hnselvtqsn ilakkpkykc dkcrksfihr sslnkhekib kgdpysngtd  
 241 qgaqsgrkhh ecadcgkfl wrtqltehqr ihtgekpfec nvcgkafrhs ssllgghenah  
 301 tgekpyqcs1 cgkafqrssss lvqhqrhhtg ekpyrcnlcg rsfrhstslt qhevthsgek  
 361 pfqckecgka fsrccslvqh erhtgekpf eccicgraflg qspsllykhmr ihkrskpyqs  
 421 nnfsalafvpn tlpqgegll tevksyhnd cgkdfghitd fsehqrlhag ensygseqtl  
 481 lgqqslshpr ekpyqcnvcg kafkrstsf iehrihtgek pyecnecgea fsrlssltqh  
 541 erhtgekpy ecidcgkafs qsssligher htgekpyec necgrfrkk tnldhqrth  
 601 tgekpyacke cgrnfsrssa ltkhhrvhar nklqes

(SEQ ID NO: 10)

Fig. 32

1 ctggcagcgg actttgaata gggaaagtttt gcaggggtta cgcttgcagt cagtccgctg  
 61 tttgcaaata ttgcgtgggc tcggcgcgt gcccggctcg ggagggtccg gacccggcgt  
 121 ccgattgcag cgccatccag tttgcatgaa actttcacct ggcgtcccg gaacagttc  
 181 tgctcggact cctgatcggtt cacccctgt tttcccgac agcgaggact gtctttcca  
 241 acccgacatg gatgtgtcc caatgttagt catcttccag gaactacaga ttgtgcacga  
 301 aacgggctac ttctcggttc tgccgtccct ggaggaatat tggcaacaga cctgcctgga  
 361 gttgaacgc tatcttcaga gtgaggccctg ctacgtgtca gcctctgaga taaaatttga  
 421 cagccaggaa gacctgtggc ccaaattcat tctagctcgg gagaagaagg aggaatcaga  
 481 actgaagatt tttcttagtc ccccaagagga ctctctgatc agctccagct ttaattataa  
 541 ctttagagacc aatagcctga actctgtatc cagcagttagt tcttcggaca gttcagagga  
 601 actttcaccc acgaccaaata ttacctctga tcccatgggt gaagtcttag tcaattcagg  
 661 aaatctgagt tcctccgtca tttccacacc tccatcttct ccagaagtga acagggaaatc  
 721 ttctcaacta tggggctgtg ggccaggaga cctgcctca cctggaaagg ttcaagtgg  
 781 gacctctggg aagtctgggt acaaggtaa tggcgacgco tccccagatg gcagaagacg  
 841 ggtacatcggt tgccacttta atggctcgag gaaagttac acgaaaagct cccacttgg  
 901 agcacatcag cgcactcaca caggagaaaa gccttacaga tgctcatggg aaggttgtga  
 961 gtggcgttt gcaagaagtg atgagttgac cagacacttc cgaaagcata ccgggtccaa  
 1021 gcctttaaa tgctccact gtgacagggt tttctccagg tctgaccacc tggccctgca  
 1081 catgaagagg cacctctgaa ggagcagagg gacgaatctt gttaggctaaa agaggcttcc  
 1141 aggctaagag gcccggatgg aaggagggat gcctgtaaaca gccaaagcat gccatggc  
 1201 ttcctatcca gttacctcca gggcccttc tttggaggt ctttgaggg ctacaaaagt  
 1261 catgtcagga gtggcatagc acccatggtg catgggttt gggtagcccc ggactcacca  
 1321 ctgggtccta accttctgag aggctctaag ctttggccg tgacgtcg cactgagaat  
 1381 gttagtggtt gggatgggtt tggtaggat ctattactga ctgtatgggtt aggcagactt  
 1441 ttttttctc cccctatgtg gtatcaaata actcgccgtc gcaacttta agaaatagaa  
 1501 atggcttcca aaagagctct ggtcatcctg gccaaaggag cagtcgacgc ggccgc

(SEQ ID NO: 11)

Fig. 33

1 ggatgagaca gaaggataga gaggaggaga gagagagaga gaagagaagc aaccagaaaat  
61 aggacgccaa taaaaaggag ccgcacttat ctgaagcctc aaggggcctg agccaggtcc  
121 ctgtttatg gcagttatga aaaattacct cctcccgatc ctggtgctc ccctggccta  
181 ctactactat tctacaaatg aagagtttag accagaaaatg ctccaggaa agaaagtgtat  
241 tgtcaactggg gccagcaaag ggatttggaa agaaaatggca tatcatctgt caaaaatggg  
301 agccatgtg gtatttactg ccaggtcgga ggaaggctc cagaaggtag tgctcgctg  
361 ccttgaactc ggagcagccct ctgctacta cattgctggc actatggaa acatgacatt  
421 tgccggagcaa tttattgtca aggccggaaa gctcatggc ggactggaca tgcttattct  
481 aaaccacatc actcagaccccgctgtctcttccatgac gacatccact ctgtgcgaag  
541 agtcatggag gtcaacttcc tcagctacgt ggtcatgagc acagccgcct tgcccatgtct  
601 gaagcagagc aatggcagca ttggcgatcat ctcccttgc gctggaaaaa tgacccagcc  
661 tatgatttgc ccctactctg caagcaagtt tgctctggat gggttctttt ccaccattag  
721 aacagaactc tacataacca aggtcaacgt gtccatcaact ctctgtgtcc ttggcctcat  
781 agacacagaa acagctatga aggaaatctc tggataatt gacgccttag ctctcccaa  
841 ggaggagtgc gccctggaga tcatcaaagg cacagctcta cgcaaaagcg aggtgtacta  
901 tgacaaatttgc ctttgcactc caatcctgttgc tggaaaccca ggaaggaaga tcatggaaatt  
961 ttttcatta cgatattata ataaggacat gttttaagt aacttaggaac tcctgagccc  
1021 tggtagtgg tcttagaaca gtcctgcctc atacttcgt aagccctacc cacaaaagta  
1081 tcttcaga gatacacaataa ttttgggtatcacatca tgagaaatttgcacact  
1141 tgcacagtga aaatgttaatt gtaataaaatg tcacaaacca ctttggccct gcagttgtga  
1201 acttgattgt aactatggat ataaacacat agtggttgtatc cggctttac ctcacactga  
1261 atgaaacaat gataactaat gtaacattaa atataataaa ggttaatataactcgtaaa  
1321 tgcaaaaaaaaaaaaaaaaaaaaaaaaaaaaa

(SEQ ID NO: 12)

Fig. 34

1 mavmknyllp ilvlflayyy ystneefrpe mlqgkkvivt gaskgigrem ayhlskmgah  
 61 vvltarseeg lqkvvsrcl e lgaasahyia gtmedmtfae qfivkagklm gglldmlilnh  
 121 itqtstslfh dihsrrvm evnflsyvvm staalpmlkq sngsiaviss lagkmtqpmi  
 181 apysaskfal dgffstirte lyitkvnsi tlcvlglidt etamkeisgi inaqaspkee  
 241 caleiikgta lrksevyydk spltpillgn pgrkimeffs lryynkdmfv sn

(SEQ ID NO: 13)

Fig. 35

1 gagacggacg gtggccaccc caagacgcgc cccagccgc catggcccg atcctccggg  
 61 catcctgcct tctgtccctg ctccctggccg gttttgttcc gcccggccgg ggacaagaga  
 121 agtctaagac agactgccat ggccgtatga gtggtaccat ctacgagtat ggagccctca  
 181 ccatcgatgg ggaggaatac attcccttta agcagtatgc aggcaaatat atccttttg  
 241 tcaacgtacg cagctactga ggtctgacag accaataacct tgaactgaat gcactacaag  
 301 aagaacttgg gccatttggc ttggtcattc tgggcttccc ttccaaccaa tttggcaaac  
 361 aggagccagg cgagaactcg gagatactcc ccagtctcaa gtatgttgc ccaggtgggg  
 421 gcttggcc taatttccag ctcttgaga aaggagatgt gaacggggag aaagagcaga  
 481 aattctacac ttcctgttca aactcctgccc ctcccaactgc agaactcctg ggctcacctg  
 541 gcccctctt ttgggaacccc atgaagatcc atgacatccg ctggacttt gagaagttcc  
 601 tggggggcc agatggcata ccgggttatgc gctggtacca ccggaccaca gtcagcaacg  
 661 tcaagatgga catcctgtct tacatgaggc ggcaggcagc cctgagcgc aggggaaagt  
 721 aactgtatgcc cccacccctac ccctacccccc tggccatcat gcaaggccg aggaggggct  
 781 cttcaggaag gaagccacat tcccagtcat tctacccccc ccccaagattt tcttttttat  
 841 tacataaaag acaaggctgg cacaactgtg tgcgttgc accgtggaca cgtgacaatt  
 901 gtcctcactgtgt gtgcattggc acacagccac gtatctgcct gtttggaaacc cagggatgg  
 961 ccatctgtgt ttacggcttg gcacaacacc ctcatatttt ttccatgtt ctgttccaaa  
 1021 tgagccaaa gaaacacaaa gttcttagtca caatggttct gctcaaacctt gaacatcatt  
 1081 cttggggccca gcatctccca catgcccaca ctacacacca ccagcctct tcttccttcc  
 1141 tgaaggaccc tcctgagccc ccaagcccat cccacagtgc tcctgagacc agccaagaca  
 1201 actgtgagcg cgatggccgt gtacccagg tcaggggtgg tgcgttctatg aaggaggggc  
 1261 ccgaagccct tggggccggg cttcccttgc gcccgtctgt ggtggccagcc cttatgtcat  
 1321 tcaggtttag gtcctcaggc agggacacta ccccccggcc tctggaggac atgtatccct  
 1381 ctcactctgt ccactgttat ctcaacaccc ccatctgcct agtaaaggc tttctgc

(SEQ ID NO: 14)

Fig. 36

1 marilrascl lslllagfvp pgrgqekskt dchggmsgti yeygaltidg eeyipfkqya  
 61 gkyilfvnva syugltdqyl elnalqeelg pfglvilgfp snqfgkqepg enseilpslk  
 121 yvrpgggfvp nfqlfekgdv ngekeqkfyt flknscppa ellgspgrlf wepmkihdir  
 181 wnfekflvgp dgipvnmrwyh rttvsnvkmd ilsymrrqaa lsargk

(SEQ ID NO: 15)

Fig. 37

1 ctgtaaagcc ccgcctcagc cccgccccct cgtcccggcc gccgcgggccc aagccggagc  
 61 aagctaggag gcagccggct ctgcggaggc aacatgtacc ggctcctgtc aagcgtgaca  
 121 gctccggctg cggccaccgc aggcccagcc tgggacggag ggcggcgcgg ggcgcacagg  
 181 cgaccgggccc tgccctgtct gggccttggc tgggccccggc gcctggggct cggctgggg  
 241 ctggctctcg ggcgaagct ggtggctggg ctgcggggcg ccgtccccat tcagtcccc  
 301 gcgggaccccg aggccgtccgg cactaccgag ttatcgacg agcaggcccct gagccccggg.  
 361 agcccgacaca cgcctgcgc gccagcagcc aggggcttct ccagagccat cgagagcagc  
 421 cgcgcattcgc tacaccggat caaggatgag gttggtgc cccgcattcgt gttggagtt  
 481 tctgttagatg gaaaagaagt ctggtcagaa gttttaggct atgcagacgt ggagaaccgc  
 541 gtaccctgtc agccagaaac ggtcatgaga atcgcaagca tcagcaaaag cctcaccatg  
 601 gtggctctgg ctaaactgtg ggaagcaggg aagctggatc tggaccttcc tgcagcac  
 661 tatgttcccg agttcccaga aaaagaatac gagggtgaaa aggtttctgt cacaacaaga  
 721 ttactaattt cgcatttaag tggatttcgt cattatgaaa aggacataaa gaaagtgaaa  
 781 gaagagaaaag cttataaagc cctgaagatg gtgaaaggga cccgcaccat atctgacca  
 841 gaaaagaac tgaaagaaaa gggaggcaaa aacaacgaaa agagcagc accgaaagcc  
 901 aaagtgcagc aggacagcga agccagatgc cgcagcgcga agccaggcaaa gaaaagaat  
 961 gacttgcac aaggcgaatt gtatttggaa gaaaaggatgg aaaattcaat tgaatcacta  
 1021 agatttatttta aaaaatgaccc tttatttctt aaacctggta gtcagttttt gtattcaacg  
 1081 tttggctata ctctgctggc agccatagta gaaaagagctt caggatataa atatttggat  
 1141 tatatgcaga aaattttcca tgattttggac atgctgacaa ctgtccagga gggaaaacgag  
 1201 ccagtgtttt acaacagagc aagattttac gtgtacaata aaaagaaacg tcttgcac  
 1261 acacccttacg tggataactc ctataaatgg gctgggtggc gatttctgtc cacagtgggt  
 1321 gaccccttgc aattttggaaa cgcacatgtg tatggctacc aagttggca gtttagaac  
 1381 tcaaatggaaa atctcttgc tggatatctc aagccagaaa caatggat gatgtggacc  
 1441 ccagtcccta acacagagat gtcctggat aagaggggaa aatatgcacat ggcgtgggt  
 1501 gtggtagaga agaagcaac gtacggatcc tgcaggaagc agccgcacta cgcctcacat  
 1561 actggagggtg ctgtgggtgc cagtagtgc ctgtggtcc ttctgtgaaa actggactca  
 1621 gaggccgtaa atacaaggat tcccccacga ggaataatcg tctctatcat atgcacacatg  
 1681 cagtcgtgg ggctcaatag cactgccttgc aagatcgctc tggatttgc taaagacaga  
 1741 gctgactaat cctaatggca gcacaggtcc acagtgagcc ttccattttt tggaaatgttgc  
 1801 acgttcccaa atacataaaac cctttaagga tacatttcgt tcccaaatac ataaaccctt  
 1861 taaggatatac ttgttaatag agtacagttt aatgtggaga attatgtacc tctaattgtc  
 1921 taattttgtc actgcctttt tattggacaa ttatgttccctt acactcaggaa aataaacagt  
 1981 tgtttctact tttaaaaaaa aatgtttact cttgaaataa aatcttctga t

(SEQ ID NO: 16)

Fig. 38

1 myrlssvta raaatagpaw dggrrgahrr pg1pvlg1gw agglglgl1 algak1vvgl  
61 rgavpiqspa dpeasgttel sheqaislgs phtpappaar gfsraiessg dllhrikdev  
121 gapgivvgvs vdgkevwseg lgyadvenrv pckpetvmri asisks1tmv alaklweagk  
181 ldldlpvqhy vpefpekeye gekvsvttr1 lishlsgirh yekdikkvke ekaykalkmv  
241 kgtpppdqe kelkekggkn neksdtpkak aeqdsearcr sakpgkknd feqgelylke  
301 kfensieslr lfkndplffk pgsqlflystf gytlalaive rasgykyldy mqkifhldm  
361 lttvqeenep viynrarfyv ynkkr1lvnt pyvdnsykwa gggflstvgd llkfgnamly  
421 gyqvgqfkns nenllpgylk petmvmmwtp vpntemswdk egkyamawgv vekqtygsc  
481 rkqrhyasht ggavgassvl lvlpeeldse avnnkvpprg iivsiicnmq svglnstalk  
541 ialefdkdra d

(SEQ ID NO: 17)

Fig. 39

1 aggctgnnag ccacacttgg gaaaggaagc atggcgtgcg agctgcgagc tggcggagg  
61 tggggcccgcg ggctgcagac tgcactgcgg gcccggcg tggctggagt tcggcggagg  
121 aagccagttc tgcacccca gaaactaca gtccagttt gggggccac acaaagtctg  
181 gcttcaggga tctctgcagg acagttatac agcacacagg cagccgagga caaggaggag  
241 gagagcctgc actccatcat cagcaacact gaggcagtgc ggggttctgt ctccaaacat  
301 gagttccagg cagagacaaa gaaactttt gacatcgtag cccgttctct gtactcagaa  
361 aaagagggtgt tcatacgaga gctcatctcc aatgccagt atgccttggaa gaaactgcgg  
421 cacaagctgg tgcgtgaagg ccaggtgctg ccagaaatgg agattcacct tcagacggat  
481 gccaagaagg gcactattac cattcaggac actggcattt ggtatgcaca ggaggagctg  
541 gtgtccaaacc ttggcacaat tgccagatcg gggtaaaagg ctttcttggaa agcactgcag  
601 aaccaggcag agaccagcag caagatcatt ggtcagttt gagtgggtt ctattcagcc  
661 ttcatggtag ctgacaaggt tgaagtctat tctcgatcag cagctccaga gagcccgagg  
721 taccagtggc ttcagatgg ttctggagtg tttgaaattt ccgaagcttc aggagttaga  
781 cctgggacca aaataatcat ccacctcaag tcaagactgta aagatttgc cagcgagtcc  
841 cgggtacaag atgtggtaac aaagtacagt aacttgc tgcctccctt gtaccttaat  
901 gggaaagcggg ttaacacttt gcaggccatc tggatgatgg acccaaagga catcagtgaa  
961 ttcatcgatg aggaattcta ccgttatatt gtcaggcctt atgataagcc cgccttcaact  
1021 ttgcactaca agacggacgc accactcaac atccgcagca tcttctatgt gccagagatg  
1081 aaaccatcca tggatgtgt gagcaggggag ctgggctcca gcgtggact gtatagccgc  
1141 aaggctctca tccagaccaa ggctgcagac atcctgccc agtggctgcg cttcattoga  
1201 ggtgtggtag atagtggaga cattccccctg aacctcagca gagagctcct gcaggagagt  
1261 ggcgtcatcc gggaaactccg ggatgttcta caacagagat tgatcaagtt cttcatttgac  
1321 cagagtaaaa aagatgtga aaaatacgca aagtttttg aagattatgg cttgttcatg  
1381 aggaggggca ttgtgaccac tgcagagcaa gacatcaagg aggatattgc aaaactgcta  
1441 cgttatgagt ctcagccct gctgtctgg cagctgacca gcttaccaga ctatgccagc  
1501 cgaatgcagg ctggcacccg caacatctat tacctgtgtg cccctaaccg tcacctggct  
1561 gaacattcac ctttacgaa agccatgaaag cagaaacata ctgaggtct cttctgtctat  
1621 gagcagttcg atgagcttac tctgtgcac ctggggagt ttgacaagaa gaagctcatc  
1681 tctgtggaaa cagacatcg cgttgcatac tacaaggagg aaaagtttgc ggacacatct  
1741 ccagctgtat agccgccttc ggagaaggaa acagaagatc taatggcgtg gatgagaat  
1801 gcaactagggt cccgtgtcac caatgtgaag gtgactttcc gcctagacac ccaccctgc  
1861 atggtgaccg tgctggagat gggggctgtc cggcattttc tgcgtatgc caagctggcc  
1921 aagacccagg aggaacgtgc ccaactgcta cagcccacac tggagatcaa ccccaaggc  
1981 acactgataa agaagcttgc ccagctgagg gagagcgcagc cggagctggc ccagctgctc  
2041 gtggatcaga tctatgagaa tgccatgata gcagcaggac tcgttgcata ccccccggcc  
2101 atggtcggcc gcctgaacca ccttttggtc aaggtcctgg agaaacactg acagccaaga  
2161 cactggattt agtgtcaacc caggtcttct cgggtgataa tggacccgtcc tggggaggca  
2221 ggacttaata cacaacactg gccaccaact gcttgagctc agcttttattt acttcaatta  
2281 aacagtattt cttagtc

(SEQ ID NO: 18)

Fig. 40

1 acelravllw grglqtvira palagvrrgk pvlhlqkttv qfrgptqsla sgisagqlys  
61 tqaaedkeee slhsiiisnte avrgsvskhe fqaetkklld ivarslysek evfirelism  
121 asdaleklrh klvceggvlp emeihlqtda kkgtitiqdt gigmtqeelv snlgtiarsg  
181 skaflealgn qaetsskiig qfgvgfysaf mvadkvevys rsaapespgy qwlsdgsgvf  
241 eiaeasgvrp gtkiiihlks dckdfasesr vqdvvtkysn fvsfplyng krintlgaiw  
301 mmdpkdisef qheefyryia qaydkprftl hyktdaplni rsifyvpemk psmfdvsrel  
361 gssvalysrk vliqtkaadl lpkwlrifrg vvdssedipln lsrellqesa lirkldvlg  
421 qrlikffidq skkdaekyak ffedylfmr egivttaeqd ikediakllr yessalpagg  
481 ltslpdyasr mqagtrniyy lcapnrhlae hspyyeamkq khtevlfcye qfdeltllhl  
541 refdkkklis vtdivvdhy keekfedtsp aderlseket edlmawmrna lgsrvtnvkv  
601 tfrldthpam vtvlemgaar hflrmqqlak tqeeraqlq ptleinprht likklcqlre  
661 sepelaqllv dqiyanamia aglvddpram vgrlndl1vk vlekh

(SEQ ID NO: 19)

Fig. 41

(SEQ ID NO: 20)

Fig. 42

1 ttcccgcgct tctgctccgc cctccgcagc cctccacagt caccggag accagccgtg  
 61 ttaagctctc tgctctgaag ctgactgact tccatggcag cccgagaagaa agcagttctg  
 121 gggccattgg tgggagcagt ggaccagggt accagctcgac cacgtttttt ggtttcaat  
 181 tcaaaaacag ctgaacttct tagtcatcat caagtagaaaa taaaacagga attcccaaga  
 241 gaaggatggg tagaacaaga cccgaaggaa attctgcagt ctgttatga gtgtatagag  
 301 aaaacgtgtg agaaaacttgg acagctcaat attgatattt ccaacatcaa agcattgg  
 361 gtcagcaacc agagggaaac cacagtagtc tggacaagg tcaccggaga gcctcttat  
 421 aatgccgtgg tggcttgc cctaagaacc cagtctactg ttgagaacct tagtaaaaaga  
 481 attccaggaa ataataactt tgtcaagtcc aagacaggcc ttccacttag cacgtatttc  
 541 agtgcagtga aacttcgttg gtccttgac aacgtaaaa aggtccaaga ggctgttgg  
 601 gaaaatagag ctcttttgg gaccattgtat tcattgccta ttggagttt aacaggagga  
 661 atccatgggg gtgtccactg tacagatgt acaaatgca acaaggacat gcttttaac  
 721 attcattctt tggaaatgggta taaagagctc tgcaattttt ttggaaattcc aatggaaatt  
 781 cttcccaacg ttccggagttc ttctgagatc tatggcctaa tgaaagctgg ggccttggaa  
 841 ggtgtaccaa tatctgggtt ttggggggac cagtctgtc tttgggtggg acaaatgtgc  
 901 ttccaggatg gacaggccaa aaacacgtat ggaacagggt gcttcttatt gtcaacacg  
 961 ggccataagt gtgtatttt tgaacatggc cttctgacaa ccgtacata taaacttggc  
 1021 agagacaaac ctgtgtat tgcgtggaa gttccgtgg ctatagctgg tgctgtatc  
 1081 cgctggctaa gagacaaccc tggaaattt aagtccctgtt agggaaatttga aaaacttgc  
 1141 aaggaagtag gtacttctt tggctgtc tac ttcgttccag cattttcagg gttatatgc  
 1201 ccttattggg agcccagtgc aagaggatc atctgtggac tcaactcattt cacaataaa  
 1261 tgtcatatcg ctttgcgtgc actagaagct gttgtttcc aaacccgaga gattttggat  
 1321 gccatgaatc gcgactgtgg aattccactc agtcatttac aggttagatgg aggaatgacc  
 1381 agcaataaaa ttctttagtca gctacaagca gacattctgtt atatccatc agtggaaaccc  
 1441 tccatgcctg aaacaactgc actaggcgt gccatggcag ctgggggtgc agaggggggtt  
 1501 ggtgtgtgg a tgcgtggaa tgaggatgg tca gctgtca caatggagcg gttgaacct  
 1561 cagatcaatg ctgaagaaaag cgaaatccgt tactccacat ggaagaaaagc tggatgaaag  
 1621 tcaattgggtt gggtttacac tcagtc tcca gaaagtggta tcccataaaat aataccacct  
 1681 cacgatttc caagatgca gcttttaat gtgatatgaa aatctgacta ttctgtctca  
 1741 tagtataatg atgctattca tagactctga ttttttcat aagccactgg ctgcattgatc  
 1801 ctctaaagcag acctatgact tggaaataaaag aaagtgcagc agaaaagaatc ctccagaaaac  
 1861 atttaatttt ttttaacat tgacagttt gatcgggtca gtcaccttgg aggtgtac  
 1921 ctgcctccac tggcatgtat tcctacacta tttccgtttaa ggtctagggt gattttggta  
 1981 tcctgtctat tggaaatgtgc cattcagtat attcagatgc tagtggatta cacaatgttg  
 2041 aggaagaggt tggttactaac ctgttccaaa tggatggctt ctgtttttt tgcttttaac  
 2101 agctcagatg tcttcttttccat tatataattt aaggccacaa cattactggat tattcaat  
 2161 gggaaacatct aaagaatttttggataattt aatttgcata ttcttgc ttaagacatt  
 2221 tttctgtaca gtttttgc caaaattcca accttgcag gtttttaca ctgtccact  
 2281 aactaccata gttttctgtc tggcttccat aggttagaac actttttttt tctgttttt  
 2341 tttcttctt ctttttata ttttttttctt gtatgtataa catacatgcc tataatattt  
 2401 atatgtcgag agtaacccat ttataaattt aagagcacattt atattcaata agtataaga  
 2461 gggctggctc taagtggact actatgtata cag

(SEQ ID NO: 21)

Fig. 43

1 maaakkavlg plvgavdqqg sstrflvfns ktaellshhq veikqefpre gwveqdpkei  
 61 lqsvyeciek tceklgqlni disnikaigv snqrettvvw dkvtgeplyn avwldlrtq  
 121 stvenlskri pgnnnfvksk tglplstyfs avklrwlldn vkkvqeavee nralfgtids  
 181 wliwslttgg i hggvhctdvt nasrtmfn hsllewdkelc effgipmeil pnvrssseiy  
 241 glmkagaleg vpisgclgdq saalvgqmcf qdggakntyq tgcfllicntg hkcvfsehgl  
 301 lttvayklgr dkpvyyaleg svaiagavir wldrnlgii sseeiekak evgtsygcif  
 361 vpafsglyap ywepsargii cgltqftnkc hiafaaleav cfqtreilda mnrdcgipls  
 421 hlqvddggmcts nkilmqlqad ilyipvvkps mpettalga maagaagvg vwslepedls  
 481 avtmerfepq inaeeeseiry stwkkavmks igwvttqspe sgip

(SEQ ID NO: 22)

Fig. 44

1 tgtcagactc tcgatttctc ctcctactcc tcctccgagg aattctgcgc cctgttaactg  
 61 ttctgcctc ccctttaaag gttgacttgc cctacggcgc tccaccgcgc tccagtctc  
 121 ttgcgcctcc tgctcaaccc gctcctgact gccccacgccc gcgttagttcc agcagcaag  
 181 cagaagggtg caccgggaga tggagagcaa agccctgctc ctgggtgtcc tggagatgg  
 241 gctccagagt ttgaccgcct tccgaggagg ggtggccgca gcagacgcag gaagagatgg  
 301 ctcagacatc gaaagcaaat ttgccttaag gacccctgaa gacacagctg aggacacttg  
 361 tcatactcatt cctggattag cagactctgt gcttaactgc cacttcaacc acagcagca  
 421 gacccctgtg gtgatccatg gatggacgg aacgggaatg tatgagagtt ggggcccaa  
 481 acttggcc gccctgtaca agagagaacc tgactccaat gtcattgttag tagactgggt  
 541 gtatcggcc cagcaacatt atccagtgtc agctggctac accaagctgg tggaaatga  
 601 tgtggccaga ttcataact ggatggagga ggagtttaag tacccttag acaacgtcca  
 661 cctcttaggg tacagcctt gagccatgc tgctggcgta gcaggaagtc tgaccaataa  
 721 gaaggtaat agaattactg gtttggatcc agctggccct aactttgagt atgcagaagc  
 781 ccccaactc ctttctcctt atgacgctga tttttagat gtcttacaca catttaccag  
 841 ggggtcacct ggtcgaagta ttggatcca gaaaccagtg gggcatgtt acatttatcc  
 901 caatggaggc actttccagc caggatgca cattggagaa gccatccgt tgattgcaga  
 961 gagaggactc ggagacgtgg accagctgtt gaagtgtctcg catgagcgtt ccattcatct  
 1021 cttcatttgc tccctgtca atgaagaaaa ccccagcaaa gcatacaggt gcaactccaa  
 1081 ggaagccctt gagaaggcgtc tctgccttag ttgttagaaag aatcgctgt acaatctgg  
 1141 ctagatgatc aacaagggtca gagccaaagag aagcagcaag atgtacccatg agactcgctc  
 1201 tcagatgccc tacaaagtgt tccattacca agtcaagatt cacttttctg ggactgagaa  
 1261 tggcaagcaaa cacaaccagg ccttgcataat ttctctgtac ggcacagtgg ccgagagcga  
 1321 gaacattttcc ttcacccctgc ccggaggttt cacaataaaa acctactct tcttgattta  
 1381 cacggagggtg gacatcgagg aactgtctcat gatcgaatgtt aagtggatga ggcactccct  
 1441 cttcactgtgg cccgactgtt ggagccccc cagcttcgtc atcgagagga tccgagtggaa  
 1501 agccggagag actcagaaaaa aggtcatctt ctgtgttagg gagaagttt ctcatctgca  
 1561 gaaggaaaag gactcagcag tttttgtt gatccatgac aagtctgtt agaagtctgg  
 1621 ctgacactgg acaaacaac aagagaagaa agcatccgag ttctttgaag acagaagaaa  
 1681 acaaagtaaa tttaatttaa aaaaataata cccttggatggatggatggatggatggatggatgg  
 1741 cctgagtatt aatcccagct ctatcttgc agttaaacag aagacagctt caaatattaa  
 1801 acggggctt acccagggtt aggaatctaa tgccccatag caggtcttcc agcatcagaa  
 1861 gacatcaggc aggagaaaaca tgctgttttgc tatcccttaa gaaggaatca tttgtccca

Fig. 45A

(SEQ ID NO: 23)

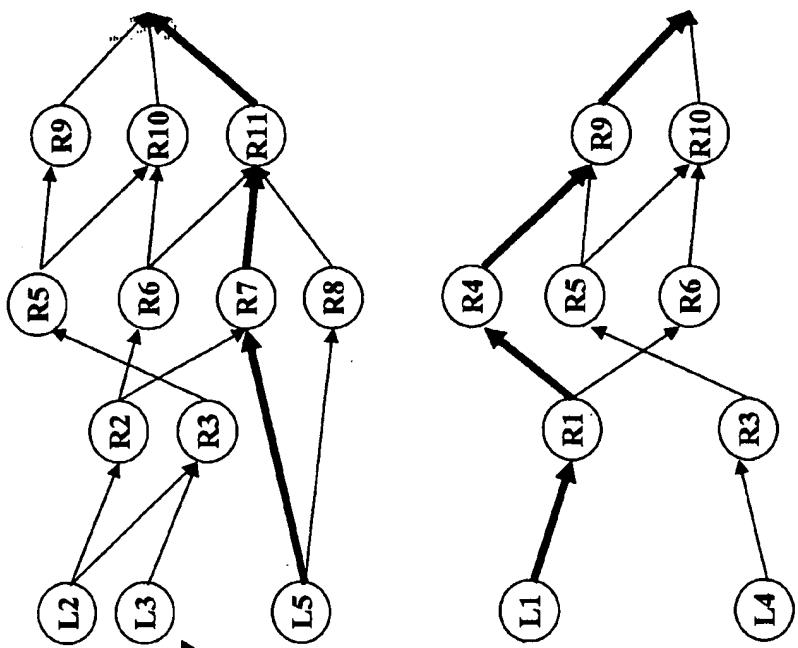
Fig. 45B

1 meskalllvv lgvwlqslta frggvaaada grdfsdiesk falrtpedta edtchlipgl  
61 adsvsnchfn hssktfvvih gwtvtgmyes wvplvaaly krepdsnviv vdwlryaqqh  
121 ypvsaqytkl vgndvarfin wmeefnypl dnvllgysl gahaagvags ltnkkvnrit  
181 gldpagpnfe yaeapsrlsp ddadfvdlh tfrtrgspgrs igiqkpvghv diypnggtfq  
241 pgcnigeair viaerglgdv dqlvkcsheh sihlfidsll neenpskayr cnskeafekg  
301 lclscknrc nnlgyeinkv rkrsskmyl ktrsqmpykv fhyqvkihfs gtengkqhng  
361 afeislygtv aesenipftl pevstnktys fliytevdig ellmmklkwm sdsyfswpdw  
421 wsspsfvier irvkagetqk kvifcarekv shlkqgkdsa vfvkchdksl kksg

(SEQ ID NO: 24)

Fig. 46

Identification of disease subtypes allows for identification of causal targets for each subtype



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graph TD
    OFP[OFP] --> Subtype1[Subtype 1]
    OFP --> Subtype2[Subtype 2]
  
```

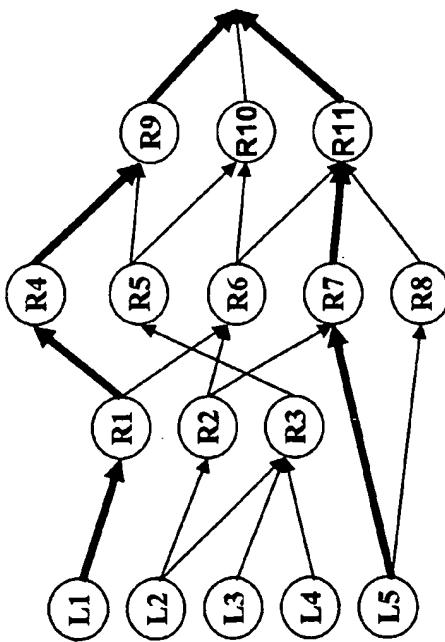


Fig. 47

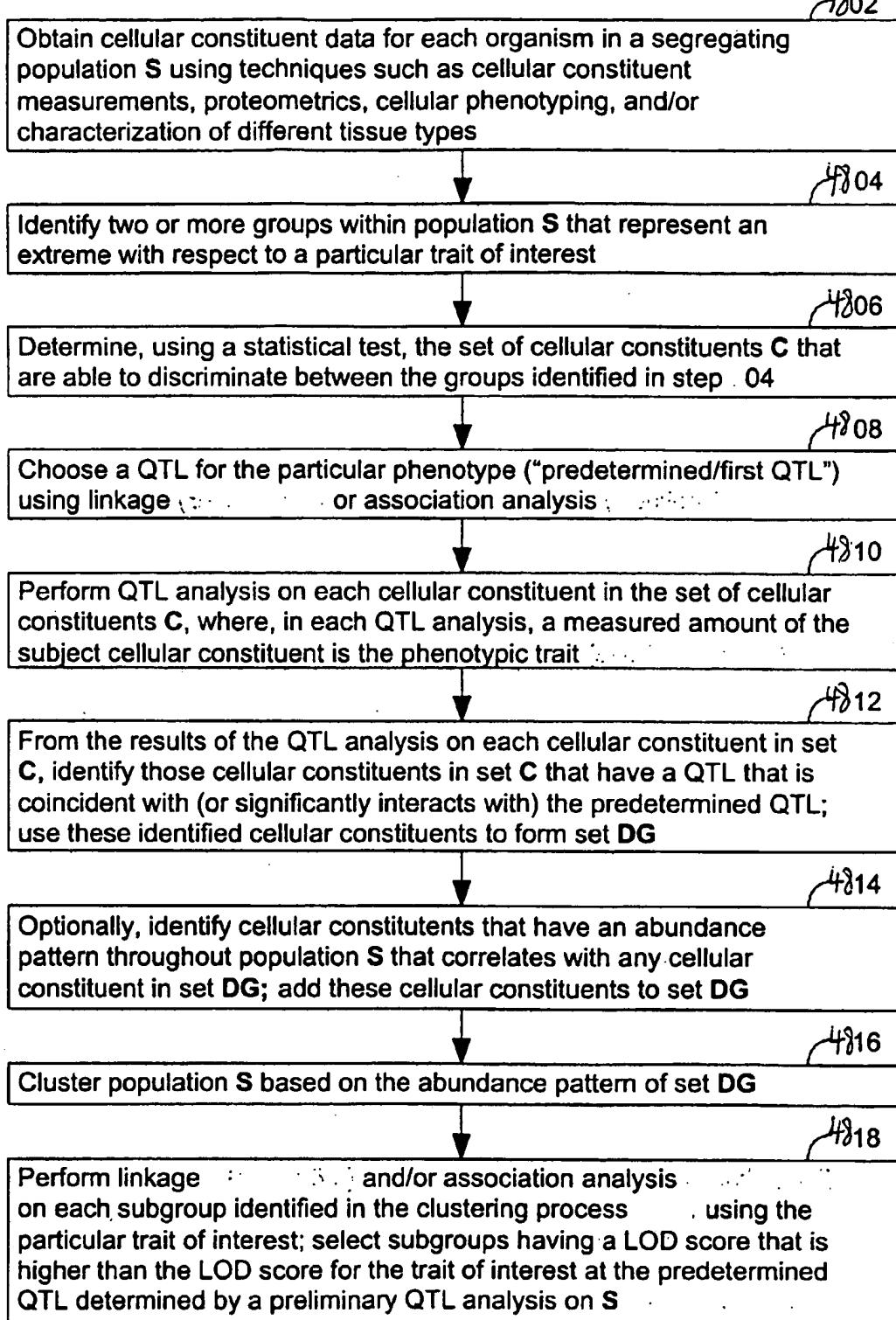


FIG. 48

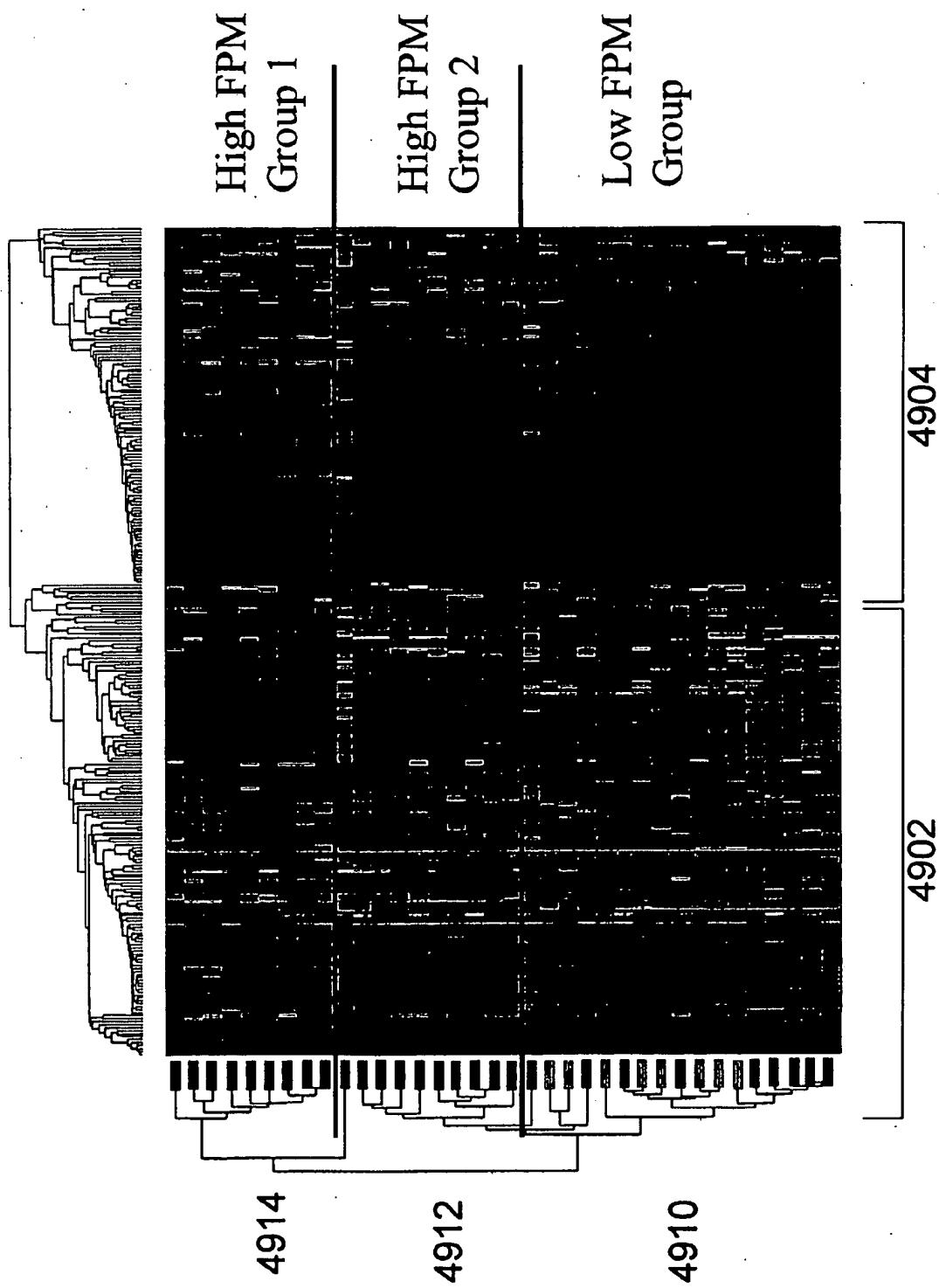
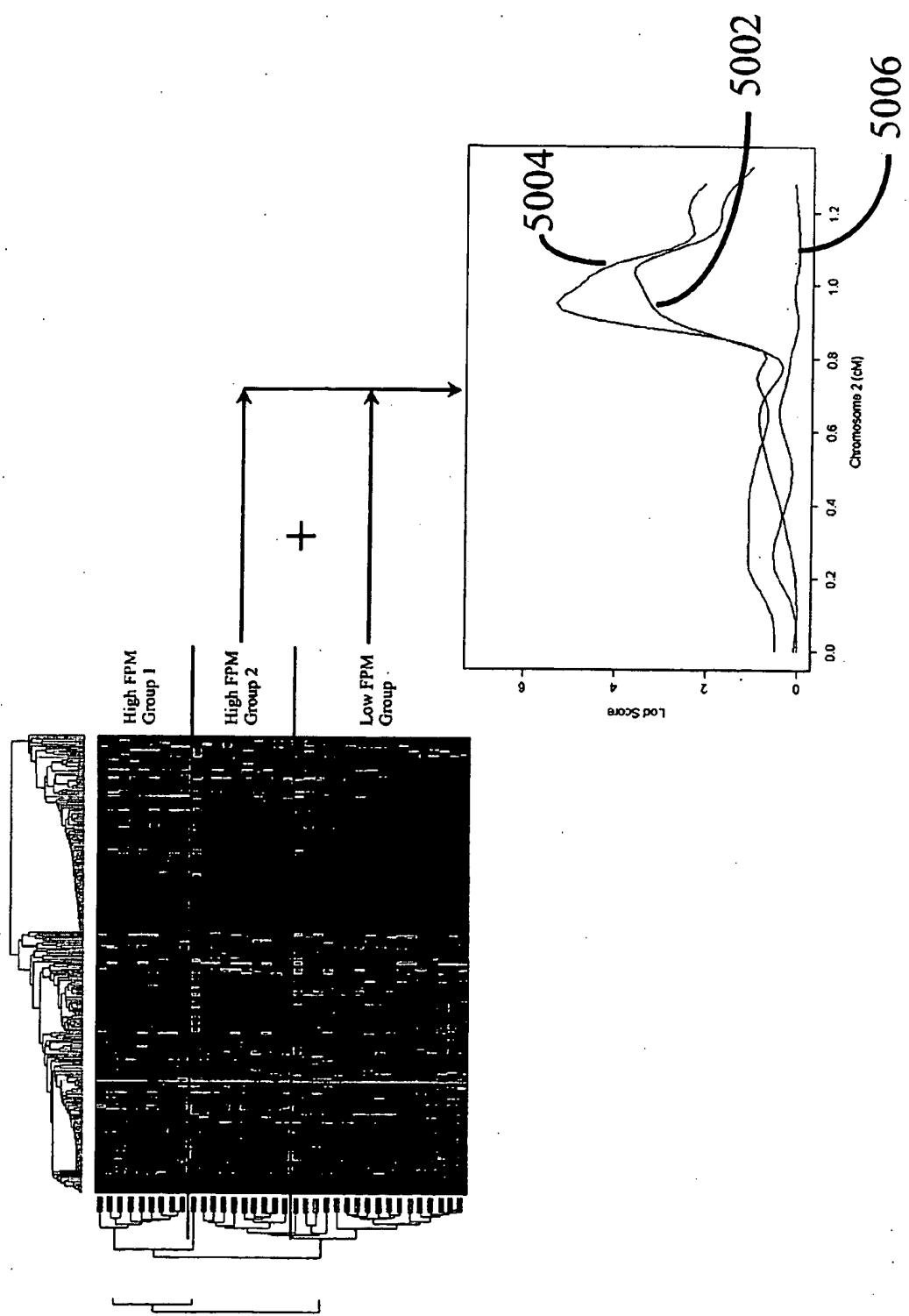


Fig. 49



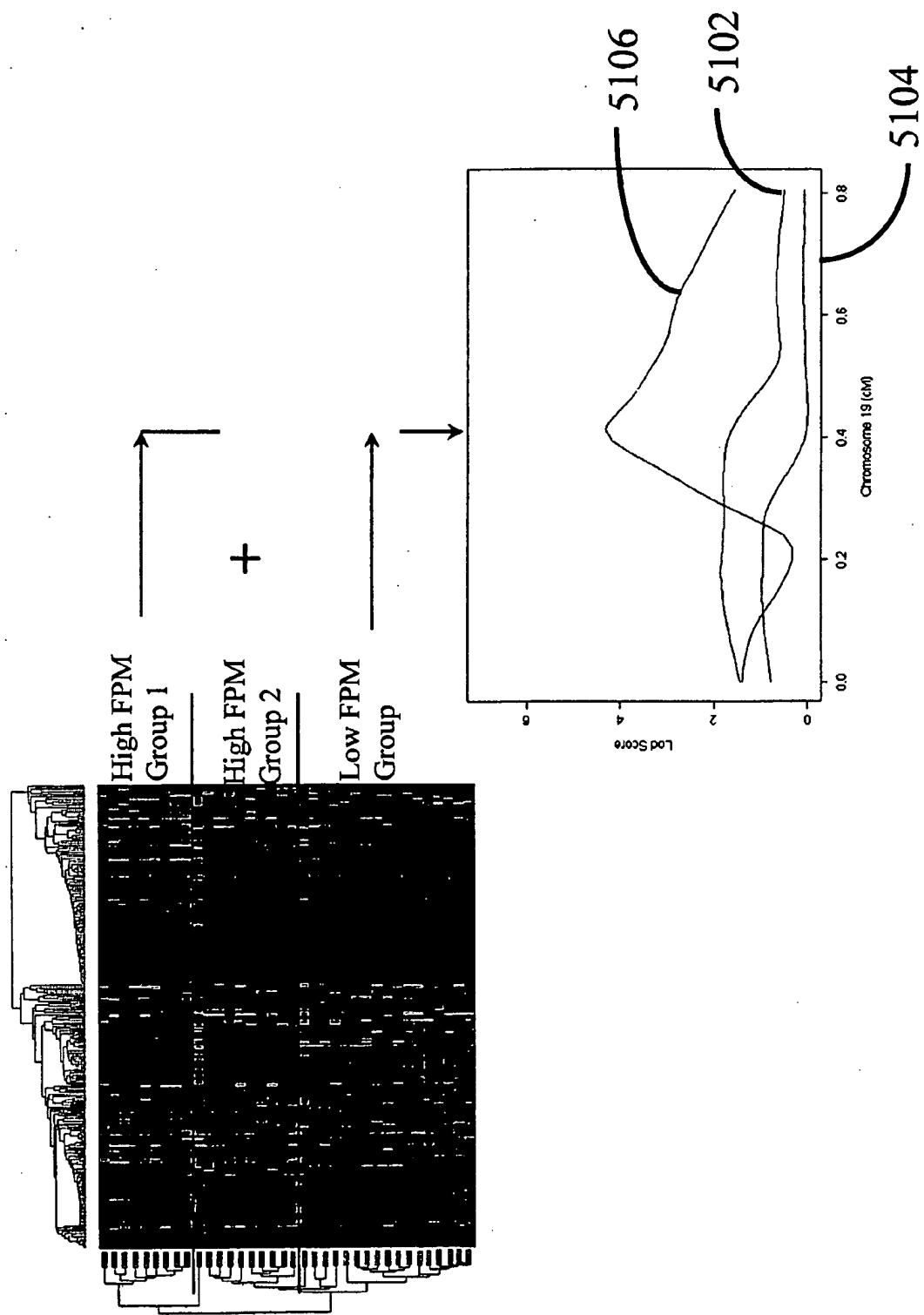


Fig. 51

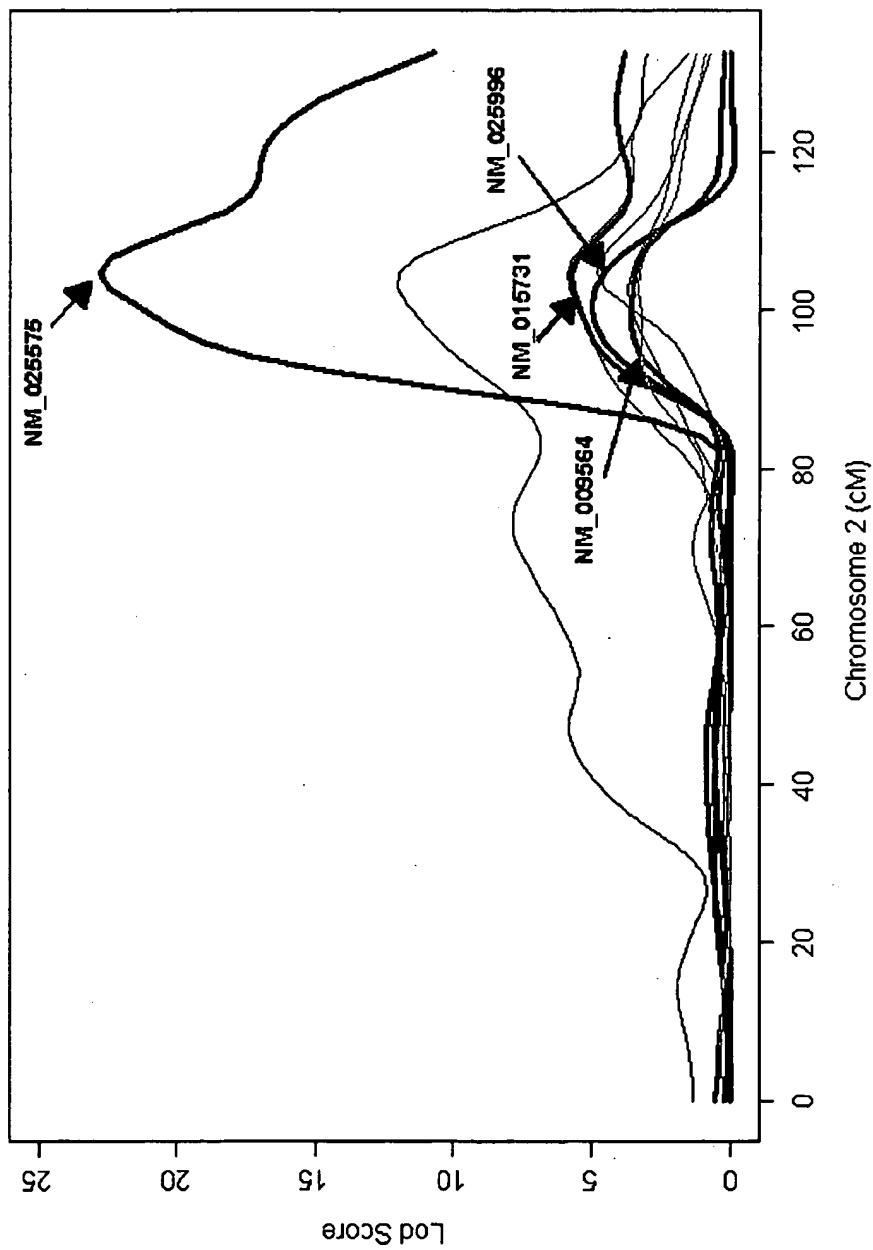


Fig. 52

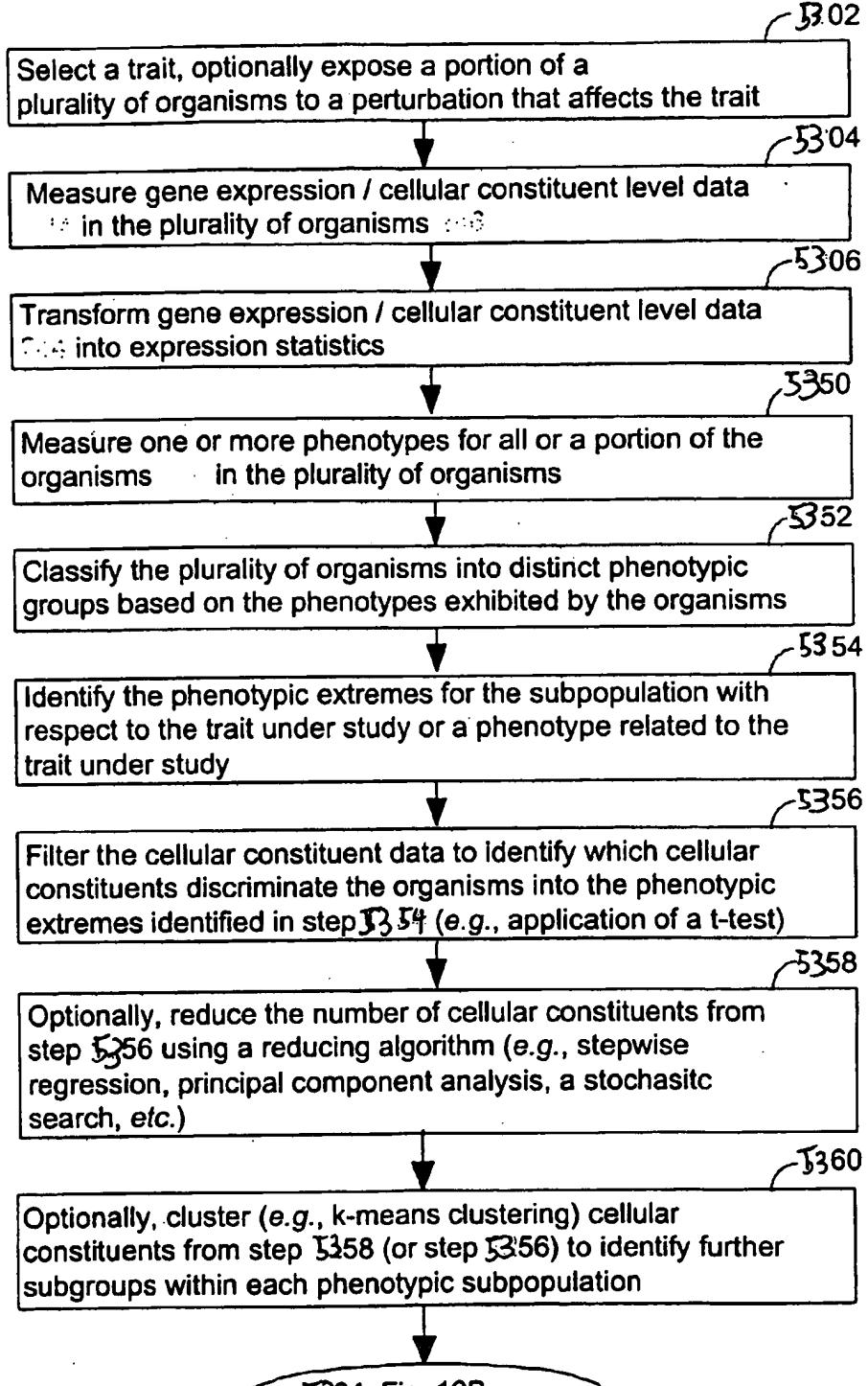
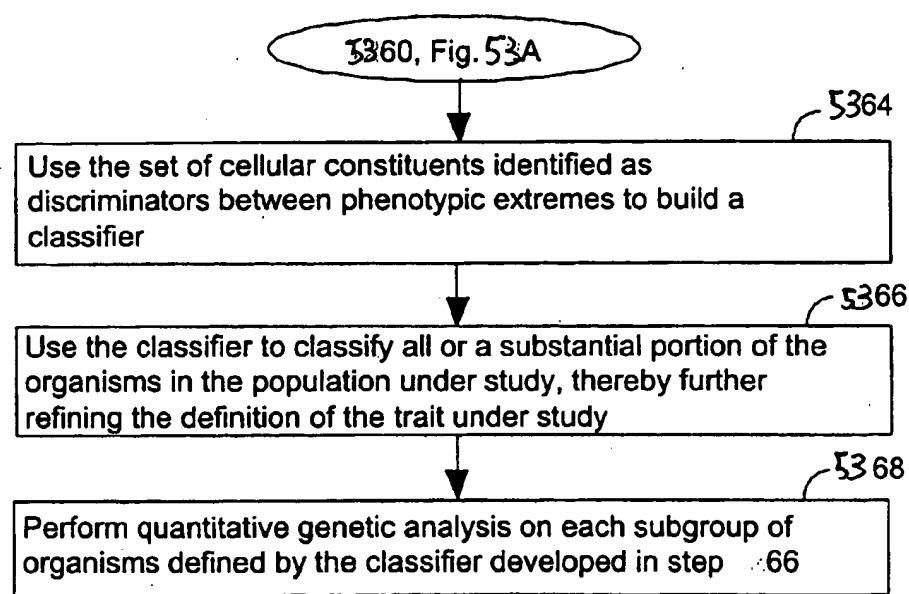
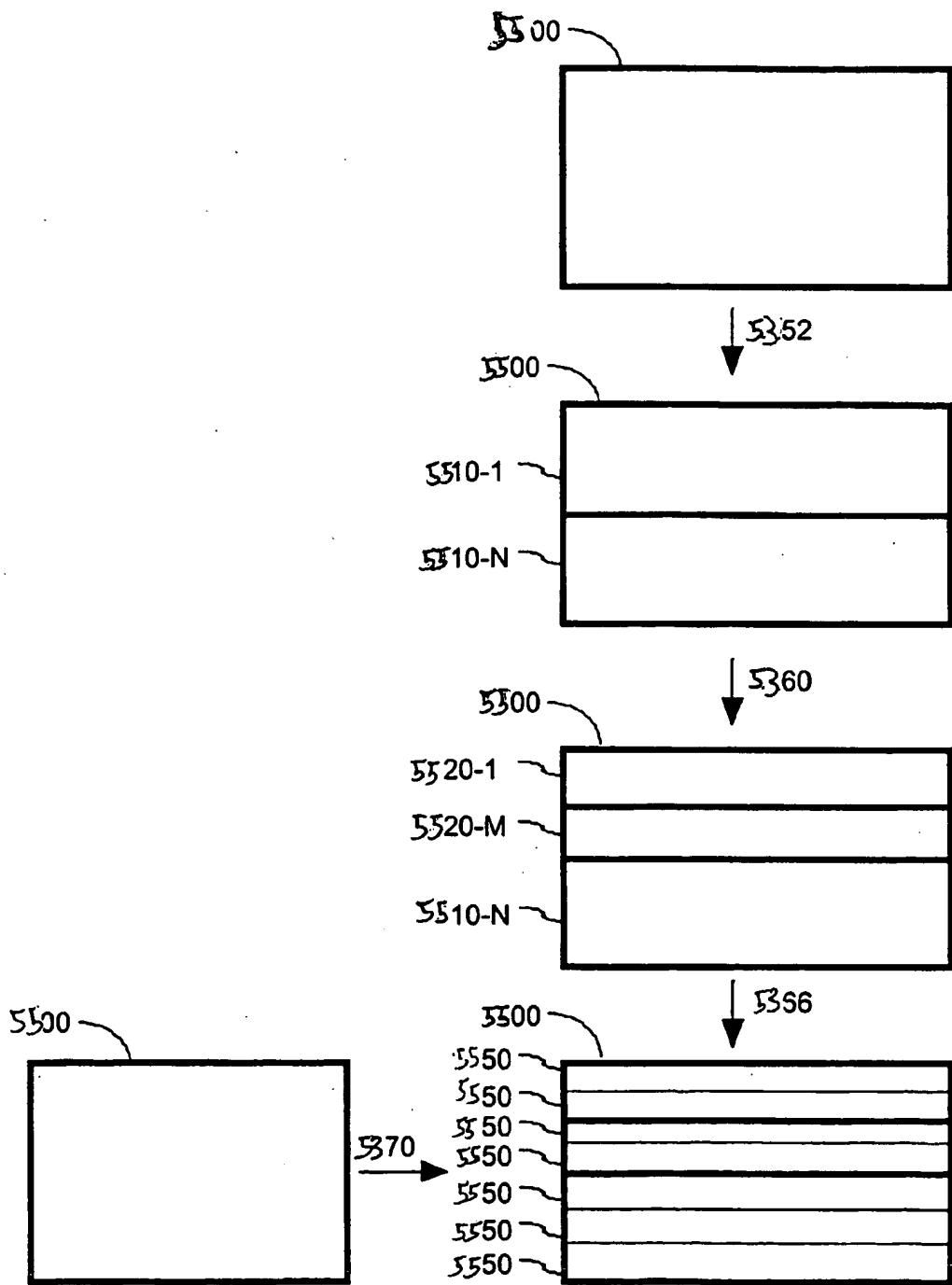


FIG. 53A

**FIG. 53B**

|               | Phenotype<br>1     | ... | Phenotype<br>M     | CC<br>248-1      | ... | CC<br>248-Z      |
|---------------|--------------------|-----|--------------------|------------------|-----|------------------|
| Organism 46-1 | Amount<br>1301-1-1 | ... | Amount<br>1301-1-M | Level<br>250-1-1 | ... | Level<br>250-1-Z |
| Organism 46-2 | Amount<br>1301-2-1 | ... | Amount<br>1301-2-M | Level<br>250-2-1 | ... | Level<br>250-2-Z |
| ⋮             | ⋮                  | ⋮   | ⋮                  | ⋮                | ⋮   | ⋮                |
| Organism 46-N | Amount<br>1301-N-1 | ... | Amount<br>1301-N-M | Level<br>250-N-1 | ... | Level<br>250-N-Z |

FIG. 54

**FIG. 55**

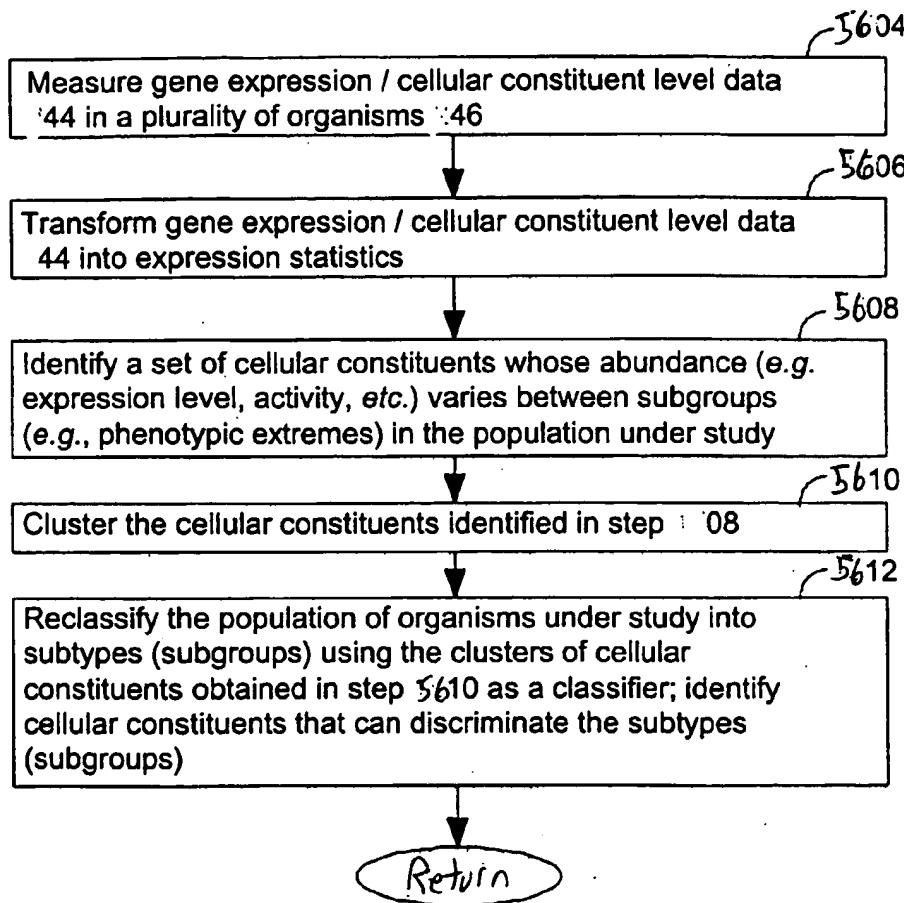


FIG. 56